

SEQUENCE LISTING

<110> Hoffmann-La Roche Inc.

<120> RA antigenic peptides

<130> 21796

<160> 141

<170> PatentIn version 3.2

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<213> Homo sapiens

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<213> Homo sapiens

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Gly	Asp	Arg	Gly	Met	Gln	Leu	Met	His	Ala	Asn	Ala	Gln	Arg	Thr	Asp
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<213> Homo sapiens

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Ile Asn Asn Gln Leu Thr Leu Asp Ser Asn Thr Lys Tyr Phe His Lys

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Ile Asn Asn Gln Leu Thr Leu Asp Ser Asn Thr Lys Tyr Phe His Lys
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Leu

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Met Pro Lys Asn Val Val Phe Val Ile Asp Lys Ser Gly Ser Met Ser
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Gly Arg Lys

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Met Pro Lys Asn Val Val Phe Val Ile Asp Lys Ser Gly Ser Met Ser
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Gly Arg

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Met Pro Lys Asn Val Val Phe Val Ile Asp Lys Ser Gly Ser Met Ser
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Gly

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Asn Val Val Phe Val Ile Asp Lys Ser Gly Ser Met Ser Gly
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Lys Asn Val Val Phe Val Ile Asp Lys Ser Gly Ser Met Ser Gly
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Asn Val Val Phe Val Ile Asp Lys Ser Gly Ser Met Ser Gly Arg
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Asn Val Val Phe Val Ile Asp Lys Ser Gly Ser Met Ser Gly Arg Lys
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Gly His Pro Gln Tyr Leu Leu Asp Ser Asn Ser Trp Ile Glu Glu
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Gly His Pro Gln Tyr Leu Leu Asp Ser Asn Ser Trp Ile Glu Glu Met
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Gly His Pro Gln Tyr Leu Leu Asp Ser Asn Ser Trp Ile Glu
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Gly His Pro Gln Tyr Leu Leu Asp Ser Asn Ser Trp Ile Glu Glu Met
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Pro Ser

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His Pro Gln Tyr Leu Leu Asp Ser Asn Ser Trp Ile Glu
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Gly His Pro Gln Tyr Leu Leu Asp Ser Asn Ser Trp
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Gly Val Asp Arg Tyr Ile Ser Lys Tyr Glu Leu Asp Lys Ala Phe Ser
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Asp Arg Asn

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Arg Tyr Ile Ser Lys Tyr Glu Leu Asp Lys Ala Phe Ser Asp Arg
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Ile Ser Lys Tyr Glu Leu Asp Lys Ala Phe Ser Asp Arg Asn
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Ile Ser Lys Tyr Glu Leu Asp Lys Ala Phe Ser Asp Arg Asn Thr
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Ile Ser Lys Tyr Glu Leu Asp Lys Ala Phe Ser Asp Arg
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Gly Ser Arg Glu Ile Lys Ser Gln Gln Ser Glu Val Thr Arg Ile Leu
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Arg Glu Ile Lys Ser Gln Gln Ser Glu Val Thr Arg
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Gly Ser Arg Glu Ile Lys Ser Gln Gln Ser Glu Val Thr Arg
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Arg Glu Ile Lys Ser Gln Gln Ser Glu Val Thr Arg Ile Leu
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Gly	Pro	His	Asp	Val	His	Val	Gln	Ile	Glu	Thr	Ser	Pro	Pro	Ala	Arg
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Asn

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Gly	Pro	His	Asp	Val	His	Val	Gln	Ile	Glu	Thr	Ser	Pro	Pro	Ala	Arg
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Asn Leu Lys

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Thr	Pro	His	Gly	Ile	Ile	Leu	Asp	Ser	Val	Asp	Ala	Ala	Phe	Ile	Cys
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Pro Gly

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Thr Pro His Gly Ile Ile Leu Asp Ser Val Asp Ala Ala Phe Ile Cys
1 5 10 15

Pro

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<400> 33

Thr Pro His Gly Ile Ile Leu Asp Ser Val Asp Ala Ala
1 5 10

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Gly Thr Pro His Gly Ile Ile Leu Asp Ser Val Asp Ala Ala
1 5 10

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<400> 35

Thr Pro His Gly Ile Ile Leu Asp Ser Val Asp Ala Ala Phe Ile
1 5 10 15

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Ile Asp Lys Glu Gly Val Ile Glu Pro Asp Thr Asp Ala Pro Gln Glu
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Met Gly

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<400> 37

Lys Glu Gly Val Ile Glu Pro Asp Thr Asp Ala Pro Gln Glu Met
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Ile Asp Lys Glu Gly Val Ile Glu Pro Asp Thr Asp Ala Pro Gln Glu
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Asp Lys Glu Gly Val Ile Glu Pro Asp Thr Asp Ala Pro Gln Glu
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<313> (1)..(261)

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Leu Leu Leu Phe Leu Pro Pro Leu Leu Leu Leu Leu Asp Val Pro Thr
20 25 30

Ala Ala Val Gln Ala Ser Pro Leu Gln Ala Leu Asp Phe Phe Gly Asn
35 40 45

Gly Pro Pro Val Asn Tyr Lys Thr Gly Asn Leu Tyr Leu Arg Gly Pro
50 55 60

Leu Lys Lys Ser Asn Ala Pro Leu Val Asn Val Thr Leu Tyr Tyr Glu
65 70 75 80

Ala Leu Cys Gly Gly Cys Arg Ala Phe Leu Ile Arg Glu Leu Phe Pro
85 90 95

Thr Trp Leu Leu Val Met Glu Ile Leu Asn Val Thr Leu Val Pro Tyr
100 105 110

Gly Asn Ala Gln Glu Gln Asn Val Ser Gly Arg Trp Glu Phe Lys Cys
115 120 125

Gln His Gly Glu Glu Glu Cys Lys Phe Asn Lys Val Glu Ala Cys Val
130 135 140

Leu Asp Glu Leu Asp Met Glu Leu Ala Phe Leu Thr Ile Val Cys Met
145 150 155 160

Glu Glu Phe Glu Asp Met Glu Arg Ser Leu Pro Leu Cys Leu Gln Leu
165 170 175

Tyr Ala Pro Gly Leu Ser Pro Asp Thr Ile Met Glu Cys Ala Met Gly
180 185 190

Asp Arg Gly Met Gln Leu Met His Ala Asn Ala Gln Arg Thr Asp Ala
195 200 205

Leu Gln Pro Pro His Glu Tyr Val Pro Trp Val Thr Val Asn Gly Lys
210 215 220

Pro Leu Glu Asp Gln Thr Gln Leu Leu Thr Leu Val Cys Gln Leu Tyr
225 230 235 240

Gln Gly Lys Lys Pro Asp Val Cys Pro Ser Ser Thr Ser Ser Leu Arg
245 250 255

Ser Val Cys Phe Lys
260

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<309> 1986-11-01
<313> (1)..(4563)

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Leu Leu Leu Leu Leu Leu Ala Gly Ala Arg Ala Glu Glu Glu Met Leu
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Glu Asn Val Ser Leu Val Cys Pro Lys Asp Ala Thr Arg Phe Lys His
35 40 45

Leu Arg Lys Tyr Thr Tyr Asn Tyr Glu Ala Glu Ser Ser Ser Gly Val
50 55 60

Pro Gly Thr Ala Asp Ser Arg Ser Ala Thr Arg Ile Asn Cys Lys Val
65 70 75 80

Glu Leu Glu Val Pro Gln Leu Cys Ser Phe Ile Leu Lys Thr Ser Gln
85 90 95

Cys Thr Leu Lys Glu Val Tyr Gly Phe Asn Pro Glu Gly Lys Ala Leu
100 105 110

Leu Lys Lys Thr Lys Asn Ser Glu Glu Phe Ala Ala Ala Met Ser Arg
115 120 125

Tyr Glu Leu Lys Leu Ala Ile Pro Glu Gly Lys Gln Val Phe Leu Tyr
130 135 140

Pro Glu Lys Asp Glu Pro Thr Tyr Ile Leu Asn Ile Lys Arg Gly Ile
145 150 155 160

Ile Ser Ala Leu Leu Val Pro Pro Glu Thr Glu Glu Ala Lys Gln Val
165 170 175

Leu Phe Leu Asp Thr Val Tyr Gly Asn Cys Ser Thr His Phe Thr Val
180 185 190

Lys Thr Arg Lys Gly Asn Val Ala Thr Glu Ile Ser Thr Glu Arg Asp
195 200 205

Leu Gly Gln Cys Asp Arg Phe Lys Pro Ile Arg Thr Gly Ile Ser Pro
210 215 220

Leu Ala Leu Ile Lys Gly Met Thr Arg Pro Leu Ser Thr Leu Ile Ser
225 230 235 240

Ser Ser Gln Ser Cys Gln Tyr Thr Leu Asp Ala Lys Arg Lys His Val
245 250 255

Ala Glu Ala Ile Cys Lys Glu Gln His Leu Phe Leu Pro Phe Ser Tyr
260 265 270

Asn Asn Lys Tyr Gly Met Val Ala Gln Val Thr Gln Thr Leu Lys Leu
275 280 285

Glu Asp Thr Pro Lys Ile Asn Ser Arg Phe Phe Gly Glu Gly Thr Lys
290 295 300

Lys Met Gly Leu Ala Phe Glu Ser Thr Lys Ser Thr Ser Pro Pro Lys
305 310 315 320

Gln Ala Glu Ala Val Leu Lys Thr Leu Gln Glu Leu Lys Lys Leu Thr
325 330 335

Ile Ser Glu Gln Asn Ile Gln Arg Ala Asn Leu Phe Asn Lys Leu Val
340 345 350

Thr Glu Leu Arg Gly Leu Ser Asp Glu Ala Val Thr Ser Leu Leu Pro
355 360 365

Gln Leu Ile Glu Val Ser Ser Pro Ile Thr Leu Gln Ala Leu Val Gln
370 375 380

Cys Gly Gln Pro Gln Cys Ser Thr His Ile Leu Gln Trp Leu Lys Arg
385 390 395 400

Val His Ala Asn Pro Leu Leu Ile Asp Val Val Thr Tyr Leu Val Ala
405 410 415

Leu Ile Pro Glu Pro Ser Ala Gln Gln Leu Arg Glu Ile Phe Asn Met
420 425 430

Ala Arg Asp Gln Arg Ser Arg Ala Thr Leu Tyr Ala Leu Ser His Ala
435 440 445

Val Asn Asn Tyr His Lys Thr Asn Pro Thr Gly Thr Gln Glu Leu Leu
450 455 460

Asp Ile Ala Asn Tyr Leu Met Glu Gln Ile Gln Asp Asp Cys Thr Gly
465 470 475 480

Asp Glu Asp Tyr Thr Tyr Leu Ile Leu Arg Val Ile Gly Asn Met Gly
485 490 495

Gln Thr Met Glu Gln Leu Thr Pro Glu Leu Lys Ser Ser Ile Leu Lys
500 505 510

Cys Val Gln Ser Thr Lys Pro Ser Leu Met Ile Gln Lys Ala Ala Ile
515 520 525

Gln Ala Leu Arg Lys Met Glu Pro Lys Asp Lys Asp Gln Glu Val Leu
530 535 540

Leu Gln Thr Phe Leu Asp Asp Ala Ser Pro Gly Asp Lys Arg Leu Ala
545 550 555 560

Ala Tyr Leu Met Leu Met Arg Ser Pro Ser Gln Ala Asp Ile Asn Lys
565 570 575

Ile Val Gln Ile Leu Pro Trp Glu Gln Asn Glu Gln Val Lys Asn Phe
580 585 590

Val Ala Ser His Ile Ala Asn Ile Leu Asn Ser Glu Glu Leu Asp Ile
595 600 605

Gln Asp Leu Lys Lys Leu Val Lys Glu Ala Leu Lys Glu Ser Gln Leu
610 615 620

Pro Thr Val Met Asp Phe Arg Lys Phe Ser Arg Asn Tyr Gln Leu Tyr
625 630 635 640

Lys Ser Val Ser Leu Pro Ser Leu Asp Pro Ala Ser Ala Lys Ile Glu
645 650 655

Gly Asn Leu Ile Phe Asp Pro Asn Asn Tyr Leu Pro Lys Glu Ser Met
660 665 670

Leu Lys Thr Thr Leu Thr Ala Phe Gly Phe Ala Ser Ala Asp Leu Ile
675 680 685

Glu Ile Gly Leu Glu Gly Lys Gly Phe Glu Pro Thr Leu Glu Ala Leu
690 695 700

Phe Gly Lys Gln Gly Phe Phe Pro Asp Ser Val Asn Lys Ala Leu Tyr
705 710 715 720

Trp Val Asn Gly Gln Val Pro Asp Gly Val Ser Lys Val Leu Val Asp
725 730 735

His Phe Gly Tyr Thr Lys Asp Asp Lys His Glu Gln Asp Met Val Asn
740 745 750

Gly Ile Met Leu Ser Val Glu Lys Leu Ile Lys Asp Leu Lys Ser Lys
755 760 765

Glu Val Pro Glu Ala Arg Ala Tyr Leu Arg Ile Leu Gly Glu Glu Leu
770 775 780

Gly Phe Ala Ser Leu His Asp Leu Gln Leu Leu Gly Lys Leu Leu Leu
785 790 795 800

Met Gly Ala Arg Thr Leu Gln Gly Ile Pro Gln Met Ile Gly Glu Val
805 810 815

Ile Arg Lys Gly Ser Lys Asn Asp Phe Phe Leu His Tyr Ile Phe Met
820 825 830

Glu Asn Ala Phe Glu Leu Pro Thr Gly Ala Gly Leu Gln Leu Gln Ile
835 840 845

Ser Ser Ser Gly Val Ile Ala Pro Gly Ala Lys Ala Gly Val Lys Leu
850 855 860

Glu Val Ala Asn Met Gln Ala Glu Leu Val Ala Lys Pro Ser Val Ser

865	870	875	880
Val Glu Phe Val Thr Asn Met Gly Ile Ile Ile Pro Asp Phe Ala Arg	885	890	895
Ser Gly Val Gln Met Asn Thr Asn Phe Phe His Glu Ser Gly Leu Glu	900	905	910
Ala His Val Ala Leu Lys Ala Gly Lys Leu Lys Phe Ile Ile Pro Ser	915	920	925
Pro Lys Arg Pro Val Lys Leu Leu Ser Gly Gly Asn Thr Leu His Leu	930	935	940
Val Ser Thr Thr Lys Thr Glu Val Ile Pro Pro Leu Ile Glu Asn Arg	945	950	955
Gln Ser Trp Ser Val Cys Lys Gln Val Phe Pro Gly Leu Asn Tyr Cys	965	970	975
Thr Ser Gly Ala Tyr Ser Asn Ala Ser Ser Thr Asp Ser Ala Ser Tyr	980	985	990
Tyr Pro Leu Thr Gly Asp Thr Arg Leu Glu Leu Glu Leu Arg Pro Thr	995	1000	1005
Gly Glu Ile Glu Gln Tyr Ser Val Ser Ala Thr Tyr Glu Leu Gln	1010	1015	1020
Arg Glu Asp Arg Ala Leu Val Asp Thr Leu Lys Phe Val Thr Gln	1025	1030	1035
Ala Glu Gly Ala Lys Gln Thr Glu Ala Thr Met Thr Phe Lys Tyr	1040	1045	1050
Asn Arg Gln Ser Met Thr Leu Ser Ser Glu Val Gln Ile Pro Asp	1055	1060	1065
Phe Asp Val Asp Leu Gly Thr Ile Leu Arg Val Asn Asp Glu Ser	1070	1075	1080
Thr Glu Gly Lys Thr Ser Tyr Arg Leu Thr Leu Asp Ile Gln Asn	1085	1090	1095

Lys	Lys	Ile	Thr	Glu	Val	Ala	Leu	Met	Gly	His	Leu	Ser	Cys	Asp
1100						1105					1110			
Thr	Lys	Glu	Glu	Arg	Lys	Ile	Lys	Gly	Val	Ile	Ser	Ile	Pro	Arg
1115						1120					1125			
Leu	Gln	Ala	Glu	Ala	Arg	Ser	Glu	Ile	Leu	Ala	His	Trp	Ser	Pro
1130						1135					1140			
Ala	Lys	Leu	Leu	Leu	Gln	Met	Asp	Ser	Ser	Ala	Thr	Ala	Tyr	Gly
1145						1150					1155			
Ser	Thr	Val	Ser	Lys	Arg	Val	Ala	Trp	His	Tyr	Asp	Glu	Glu	Lys
1160						1165					1170			
Ile	Glu	Phe	Glu	Trp	Asn	Thr	Gly	Thr	Asn	Val	Asp	Thr	Lys	Lys
1175						1180					1185			
Met	Thr	Ser	Asn	Phe	Pro	Val	Asp	Leu	Ser	Asp	Tyr	Pro	Lys	Ser
1190						1195					1200			
Leu	His	Met	Tyr	Ala	Asn	Arg	Leu	Leu	Asp	His	Arg	Val	Pro	Glu
1205						1210					1215			
Thr	Asp	Met	Thr	Phe	Arg	His	Val	Gly	Ser	Lys	Leu	Ile	Val	Ala
1220						1225					1230			
Met	Ser	Ser	Trp	Leu	Gln	Lys	Ala	Ser	Gly	Ser	Leu	Pro	Tyr	Thr
1235						1240					1245			
Gln	Thr	Leu	Gln	Asp	His	Leu	Asn	Ser	Leu	Lys	Glu	Phe	Asn	Leu
1250						1255					1260			
Gln	Asn	Met	Gly	Leu	Pro	Asp	Phe	His	Ile	Pro	Glu	Asn	Leu	Phe
1265						1270					1275			
Leu	Lys	Ser	Asp	Gly	Arg	Val	Lys	Tyr	Thr	Leu	Asn	Lys	Asn	Ser
1280						1285					1290			
Leu	Lys	Ile	Glu	Ile	Pro	Leu	Pro	Phe	Gly	Gly	Lys	Ser	Ser	Arg
1295						1300					1305			

Asp Leu Lys Met Leu Glu Thr Val Arg Thr Pro Ala Leu His Phe
1310 1315 1320

Lys Ser Val Gly Phe His Leu Pro Ser Arg Glu Phe Gln Val Pro
1325 1330 1335

Thr Phe Thr Ile Pro Lys Leu Tyr Gln Leu Gln Val Pro Leu Leu
1340 1345 1350

Gly Val Leu Asp Leu Ser Thr Asn Val Tyr Ser Asn Leu Tyr Asn
1355 1360 1365

Trp Ser Ala Ser Tyr Ser Gly Gly Asn Thr Ser Thr Asp His Phe
1370 1375 1380

Ser Leu Arg Ala Arg Tyr His Met Lys Ala Asp Ser Val Val Asp
1385 1390 1395

Leu Leu Ser Tyr Asn Val Gln Gly Ser Gly Glu Thr Thr Tyr Asp
1400 1405 1410

His Lys Asn Thr Phe Thr Leu Ser Cys Asp Gly Ser Leu Arg His
1415 1420 1425

Lys Phe Leu Asp Ser Asn Ile Lys Phe Ser His Val Glu Lys Leu
1430 1435 1440

Gly Asn Asn Pro Val Ser Lys Gly Leu Leu Ile Phe Asp Ala Ser
1445 1450 1455

Ser Ser Trp Gly Pro Gln Met Ser Ala Ser Val His Leu Asp Ser
1460 1465 1470

Lys Lys Lys Gln His Leu Phe Val Lys Glu Val Lys Ile Asp Gly
1475 1480 1485

Gln Phe Arg Val Ser Ser Phe Tyr Ala Lys Gly Thr Tyr Gly Leu
1490 1495 1500

Ser Cys Gln Arg Asp Pro Asn Thr Gly Arg Leu Asn Gly Glu Ser
1505 1510 1515

Asn	Leu	Arg	Phe	Asn	Ser	Ser	Tyr	Leu	Gln	Gly	Thr	Asn	Gln	Ile
1520						1525						1530		
Thr	Gly	Arg	Tyr	Glu	Asp	Gly	Thr	Leu	Ser	Leu	Thr	Ser	Thr	Ser
1535						1540						1545		
Asp	Leu	Gln	Ser	Gly	Ile	Ile	Lys	Asn	Thr	Ala	Ser	Leu	Lys	Tyr
1550						1555						1560		
Glu	Asn	Tyr	Glu	Leu	Thr	Leu	Lys	Ser	Asp	Thr	Asn	Gly	Lys	Tyr
1565						1570						1575		
Lys	Asn	Phe	Ala	Thr	Ser	Asn	Lys	Met	Asp	Met	Thr	Phe	Ser	Lys
1580						1585						1590		
Gln	Asn	Ala	Leu	Leu	Arg	Ser	Glu	Tyr	Gln	Ala	Asp	Tyr	Glu	Ser
1595						1600						1605		
Leu	Arg	Phe	Phe	Ser	Leu	Leu	Ser	Gly	Ser	Leu	Asn	Ser	His	Gly
1610						1615						1620		
Leu	Glu	Leu	Asn	Ala	Asp	Ile	Leu	Gly	Thr	Asp	Lys	Ile	Asn	Ser
1625						1630						1635		
Gly	Ala	His	Lys	Ala	Thr	Leu	Arg	Ile	Gly	Gln	Asp	Gly	Ile	Ser
1640						1645						1650		
Thr	Ser	Ala	Thr	Thr	Asn	Leu	Lys	Cys	Ser	Leu	Leu	Val	Leu	Glu
1655						1660						1665		
Asn	Glu	Leu	Asn	Ala	Glu	Leu	Gly	Leu	Ser	Gly	Ala	Ser	Met	Lys
1670						1675						1680		
Leu	Thr	Thr	Asn	Gly	Arg	Phe	Arg	Glu	His	Asn	Ala	Lys	Phe	Ser
1685						1690						1695		
Leu	Asp	Gly	Lys	Ala	Ala	Leu	Thr	Glu	Leu	Ser	Leu	Gly	Ser	Ala
1700						1705						1710		
Tyr	Gln	Ala	Met	Ile	Leu	Gly	Val	Asp	Ser	Lys	Asn	Ile	Phe	Asn
1715						1720						1725		
Phe	Lys	Val	Ser	Gln	Glu	Gly	Leu	Lys	Leu	Ser	Asn	Asp	Met	Met

1730		1735		1740
Gly Ser Tyr Ala Glu Met Lys Phe Asp His Thr Asn Ser Leu Asn				
1745		1750		1755
Ile Ala Gly Leu Ser Leu Asp Phe Ser Ser Lys Leu Asp Asn Ile				
1760		1765		1770
Tyr Ser Ser Asp Lys Phe Tyr Lys Gln Thr Val Asn Leu Gln Leu				
1775		1780		1785
Gln Pro Tyr Ser Leu Val Thr Thr Leu Asn Ser Asp Leu Lys Tyr				
1790		1795		1800
Asn Ala Leu Asp Leu Thr Asn Asn Gly Lys Leu Arg Leu Glu Pro				
1805		1810		1815
Leu Lys Leu His Val Ala Gly Asn Leu Lys Gly Ala Tyr Gln Asn				
1820		1825		1830
Asn Glu Ile Lys His Ile Tyr Ala Ile Ser Ser Ala Ala Leu Ser				
1835		1840		1845
Ala Ser Tyr Lys Ala Asp Thr Val Ala Lys Val Gln Gly Val Glu				
1850		1855		1860
Phe Ser His Arg Leu Asn Thr Asp Ile Ala Gly Leu Ala Ser Ala				
1865		1870		1875
Ile Asp Met Ser Thr Asn Tyr Asn Ser Asp Ser Leu His Phe Ser				
1880		1885		1890
Asn Val Phe Arg Ser Val Met Ala Pro Phe Thr Met Thr Ile Asp				
1895		1900		1905
Ala His Thr Asn Gly Asn Gly Lys Leu Ala Leu Trp Gly Glu His				
1910		1915		1920
Thr Gly Gln Leu Tyr Ser Lys Phe Leu Leu Lys Ala Glu Pro Leu				
1925		1930		1935
Ala Phe Thr Phe Ser His Asp Tyr Lys Gly Ser Thr Ser His His				
1940		1945		1950

Leu Val Ser Arg Lys Ser Ile Ser Ala Ala Leu Glu His Lys Val
1955 1960 1965

Ser Ala Leu Leu Thr Pro Ala Glu Gln Thr Gly Thr Trp Lys Leu
1970 1975 1980

Lys Thr Gln Phe Asn Asn Asn Glu Tyr Ser Gln Asp Leu Asp Ala
1985 1990 1995

Tyr Asn Thr Lys Asp Lys Ile Gly Val Glu Leu Thr Gly Arg Thr
2000 2005 2010

Leu Ala Asp Leu Thr Leu Leu Asp Ser Pro Ile Lys Val Pro Leu
2015 2020 2025

Leu Leu Ser Glu Pro Ile Asn Ile Ile Asp Ala Leu Glu Met Arg
2030 2035 2040

Asp Ala Val Glu Lys Pro Gln Glu Phe Thr Ile Val Ala Phe Val
2045 2050 2055

Lys Tyr Asp Lys Asn Gln Asp Val His Ser Ile Asn Leu Pro Phe
2060 2065 2070

Phe Glu Thr Leu Gln Glu Tyr Phe Glu Arg Asn Arg Gln Thr Ile
2075 2080 2085

Ile Val Val Val Glu Asn Val Gln Arg Asn Leu Lys His Ile Asn
2090 2095 2100

Ile Asp Gln Phe Val Arg Lys Tyr Arg Ala Ala Leu Gly Lys Leu
2105 2110 2115

Pro Gln Gln Ala Asn Asp Tyr Leu Asn Ser Phe Asn Trp Glu Arg
2120 2125 2130

Gln Val Ser His Ala Lys Glu Lys Leu Thr Ala Leu Thr Lys Lys
2135 2140 2145

Tyr Arg Ile Thr Glu Asn Asp Ile Gln Ile Ala Leu Asp Asp Ala
2150 2155 2160

Lys Ile	Asn Phe Asn Glu	Lys	Leu Ser Gln Leu Gln	Thr Tyr Met
2165		2170		2175
Ile Gln	Phe Asp Gln Tyr	Ile	Lys Asp Ser Tyr Asp	Leu His Asp
2180		2185		2190
Leu Lys	Ile Ala Ile Ala Asn	Ile Ile Asp Glu	Ile	Ile Glu Lys
2195		2200		2205
Leu Lys	Ser Leu Asp Glu His	Tyr His Ile Arg	Val	Asn Leu Val
2210		2215		2220
Lys Thr	Ile His Asp Leu His	Leu Phe Ile Glu	Asn	Ile Asp Phe
2225		2230		2235
Asn Lys	Ser Gly Ser Ser Thr	Ala Ser Trp Ile	Gln	Asn Val Asp
2240		2245		2250
Thr Lys	Tyr Gln Ile Arg Ile	Gln Ile Gln Glu	Lys	Leu Gln Gln
2255		2260		2265
Leu Lys	Arg His Ile Gln Asn	Ile Asp Ile Gln	His	Leu Ala Gly
2270		2275		2280
Lys Leu	Lys Gln His Ile Glu	Ala Ile Asp Val	Arg	Val Leu Leu
2285		2290		2295
Asp Gln	Leu Gly Thr Thr Ile	Ser Phe Glu Arg	Ile	Asn Asp Val
2300		2305		2310
Leu Glu	His Val Lys His Phe	Val Ile Asn Leu	Ile	Gly Asp Phe
2315		2320		2325
Glu Val	Ala Glu Lys Ile Asn	Ala Phe Arg Ala	Lys	Val His Glu
2330		2335		2340
Leu Ile	Glu Arg Tyr Glu Val	Asp Gln Gln Ile	Gln	Val Leu Met
2345		2350		2355
Asp Lys	Leu Val Glu Leu Thr	His Gln Tyr Lys	Leu	Lys Glu Thr
2360		2365		2370

Ile Gln Lys Leu Ser Asn Val Leu Gln Gln Val Lys Ile Lys Asp
2375 2380 2385

Tyr Phe Glu Lys Leu Val Gly Phe Ile Asp Asp Ala Val Lys Lys
2390 2395 2400

Leu Asn Glu Leu Ser Phe Lys Thr Phe Ile Glu Asp Val Asn Lys
2405 2410 2415

Phe Leu Asp Met Leu Ile Lys Lys Leu Lys Ser Phe Asp Tyr His
2420 2425 2430

Gln Phe Val Asp Glu Thr Asn Asp Lys Ile Arg Glu Val Thr Gln
2435 2440 2445

Arg Leu Asn Gly Glu Ile Gln Ala Leu Glu Leu Pro Gln Lys Ala
2450 2455 2460

Glu Ala Leu Lys Leu Phe Leu Glu Glu Thr Lys Ala Thr Val Ala
2465 2470 2475

Val Tyr Leu Glu Ser Leu Gln Asp Thr Lys Ile Thr Leu Ile Ile
2480 2485 2490

Asn Trp Leu Gln Glu Ala Leu Ser Ser Ala Ser Leu Ala His Met
2495 2500 2505

Lys Ala Lys Phe Arg Glu Thr Leu Glu Asp Thr Arg Asp Arg Met
2510 2515 2520

Tyr Gln Met Asp Ile Gln Gln Glu Leu Gln Arg Tyr Leu Ser Leu
2525 2530 2535

Val Gly Gln Val Tyr Ser Thr Leu Val Thr Tyr Ile Ser Asp Trp
2540 2545 2550

Trp Thr Leu Ala Ala Lys Asn Leu Thr Asp Phe Ala Glu Gln Tyr
2555 2560 2565

Ser Ile Gln Asp Trp Ala Lys Arg Met Lys Ala Leu Val Glu Gln
2570 2575 2580

Gly Phe Thr Val Pro Glu Ile Lys Thr Ile Leu Gly Thr Met Pro

2585		2590		2595
Ala Phe Glu Val Ser Leu Gln	Ala Leu Gln Lys	Ala Thr Phe Gln		
2600	2605	2610		
Thr Pro Asp Phe Ile Val Pro	Leu Thr Asp Leu Arg	Ile Pro Ser		
2615	2620	2625		
Val Gln Ile Asn Phe Lys Asp	Leu Lys Asn Ile Lys	Ile Pro Ser		
2630	2635	2640		
Arg Phe Ser Thr Pro Glu Phe	Thr Ile Leu Asn Thr	Phe His Ile		
2645	2650	2655		
Pro Ser Phe Thr Ile Asp Phe	Val Glu Met Lys Val	Lys Ile Ile		
2660	2665	2670		
Arg Thr Ile Asp Gln Met Gln	Asn Ser Glu Leu Gln	Trp Pro Val		
2675	2680	2685		
Pro Asp Ile Tyr Leu Arg Asp	Leu Lys Val Glu Asp	Ile Pro Leu		
2690	2695	2700		
Ala Arg Ile Thr Leu Pro Asp	Phe Arg Leu Pro Glu	Ile Ala Ile		
2705	2710	2715		
Pro Glu Phe Ile Ile Pro Thr	Leu Asn Leu Asn Asp	Phe Gln Val		
2720	2725	2730		
Pro Asp Leu His Ile Pro Glu	Phe Gln Leu Pro His	Ile Ser His		
2735	2740	2745		
Thr Ile Glu Val Pro Thr Phe	Gly Lys Leu Tyr Ser	Ile Leu Lys		
2750	2755	2760		
Ile Gln Ser Pro Leu Phe Thr	Leu Asp Ala Asn Ala	Asp Ile Gly		
2765	2770	2775		
Asn Gly Thr Thr Ser Ala Asn	Glu Ala Gly Ile Ala	Ala Ser Ile		
2780	2785	2790		
Thr Ala Lys Gly Glu Ser Lys	Leu Glu Val Leu Asn	Phe Asp Phe		
2795	2800	2805		

Gln	Ala	Asn	Ala	Gln	Leu	Ser	Asn	Pro	Lys	Ile	Asn	Pro	Leu	Ala
2810						2815					2820			
Leu	Lys	Glu	Ser	Val	Lys	Phe	Ser	Ser	Lys	Tyr	Leu	Arg	Thr	Glu
2825						2830					2835			
His	Gly	Ser	Glu	Met	Leu	Phe	Phe	Gly	Asn	Ala	Ile	Glu	Gly	Lys
2840						2845					2850			
Ser	Asn	Thr	Val	Ala	Ser	Leu	His	Thr	Glu	Lys	Asn	Thr	Leu	Glu
2855						2860					2865			
Leu	Ser	Asn	Gly	Val	Ile	Val	Lys	Ile	Asn	Asn	Gln	Leu	Thr	Leu
2870						2875					2880			
Asp	Ser	Asn	Thr	Lys	Tyr	Phe	His	Lys	Leu	Asn	Ile	Pro	Lys	Leu
2885						2890					2895			
Asp	Phe	Ser	Ser	Gln	Ala	Asp	Leu	Arg	Asn	Glu	Ile	Lys	Thr	Leu
2900						2905					2910			
Leu	Lys	Ala	Gly	His	Ile	Ala	Trp	Thr	Ser	Ser	Gly	Lys	Gly	Ser
2915						2920					2925			
Trp	Lys	Trp	Ala	Cys	Pro	Arg	Phe	Ser	Asp	Glu	Gly	Thr	His	Glu
2930						2935					2940			
Ser	Gln	Ile	Ser	Phe	Thr	Ile	Glu	Gly	Pro	Leu	Thr	Ser	Phe	Gly
2945						2950					2955			
Leu	Ser	Asn	Lys	Ile	Asn	Ser	Lys	His	Leu	Arg	Val	Asn	Gln	Asn
2960						2965					2970			
Leu	Val	Tyr	Glu	Ser	Gly	Ser	Leu	Asn	Phe	Ser	Lys	Leu	Glu	Ile
2975						2980					2985			
Gln	Ser	Gln	Val	Asp	Ser	Gln	His	Val	Gly	His	Ser	Val	Leu	Thr
2990						2995					3000			
Ala	Lys	Gly	Met	Ala	Leu	Phe	Gly	Glu	Gly	Lys	Ala	Glu	Phe	Thr
3005						3010					3015			

Gly	Arg	His	Asp	Ala	His	Leu	Asn	Gly	Lys	Val	Ile	Gly	Thr	Leu
3020						3025					3030			
Lys	Asn	Ser	Leu	Phe	Phe	Ser	Ala	Gln	Pro	Phe	Glu	Ile	Thr	Ala
3035						3040					3045			
Ser	Thr	Asn	Asn	Glu	Gly	Asn	Leu	Lys	Val	Arg	Phe	Pro	Leu	Arg
3050						3055					3060			
Leu	Thr	Gly	Lys	Ile	Asp	Phe	Leu	Asn	Asn	Tyr	Ala	Leu	Phe	Leu
3065						3070					3075			
Ser	Pro	Ser	Ala	Gln	Gln	Ala	Ser	Trp	Gln	Val	Ser	Ala	Arg	Phe
3080						3085					3090			
Asn	Gln	Tyr	Lys	Tyr	Asn	Gln	Asn	Phe	Ser	Ala	Gly	Asn	Asn	Glu
3095						3100					3105			
Asn	Ile	Met	Glu	Ala	His	Val	Gly	Ile	Asn	Gly	Glu	Ala	Asn	Leu
3110						3115					3120			
Asp	Phe	Leu	Asn	Ile	Pro	Leu	Thr	Ile	Pro	Glu	Met	Arg	Leu	Pro
3125						3130					3135			
Tyr	Thr	Ile	Ile	Thr	Thr	Pro	Pro	Leu	Lys	Asp	Phe	Ser	Leu	Trp
3140						3145					3150			
Glu	Lys	Thr	Gly	Leu	Lys	Glu	Phe	Leu	Lys	Thr	Thr	Lys	Gln	Ser
3155						3160					3165			
Phe	Asp	Leu	Ser	Val	Lys	Ala	Gln	Tyr	Lys	Lys	Asn	Lys	His	Arg
3170						3175					3180			
His	Ser	Ile	Thr	Asn	Pro	Leu	Ala	Val	Leu	Cys	Glu	Phe	Ile	Ser
3185						3190					3195			
Gln	Ser	Ile	Lys	Ser	Phe	Asp	Arg	His	Phe	Glu	Lys	Asn	Arg	Asn
3200						3205					3210			
Asn	Ala	Leu	Asp	Phe	Val	Thr	Lys	Ser	Tyr	Asn	Glu	Thr	Lys	Ile
3215						3220					3225			

Lys Phe Asp Lys Tyr Lys Ala Glu Lys Ser His Asp Glu Leu Pro
3230 3235 3240

Arg Thr Phe Gln Ile Pro Gly Tyr Thr Val Pro Val Val Asn Val
3245 3250 3255

Glu Val Ser Pro Phe Thr Ile Glu Met Ser Ala Phe Gly Tyr Val
3260 3265 3270

Phe Pro Lys Ala Val Ser Met Pro Ser Phe Ser Ile Leu Gly Ser
3275 3280 3285

Asp Val Arg Val Pro Ser Tyr Thr Leu Ile Leu Pro Ser Leu Glu
3290 3295 3300

Leu Pro Val Leu His Val Pro Arg Asn Leu Lys Leu Ser Leu Pro
3305 3310 3315

His Phe Lys Glu Leu Cys Thr Ile Ser His Ile Phe Ile Pro Ala
3320 3325 3330

Met Gly Asn Ile Thr Tyr Asp Phe Ser Phe Lys Ser Ser Val Ile
3335 3340 3345

Thr Leu Asn Thr Asn Ala Glu Leu Phe Asn Gln Ser Asp Ile Val
3350 3355 3360

Ala His Leu Leu Ser Ser Ser Ser Ser Val Ile Asp Ala Leu Gln
3365 3370 3375

Tyr Lys Leu Glu Gly Thr Thr Arg Leu Thr Arg Lys Arg Gly Leu
3380 3385 3390

Lys Leu Ala Thr Ala Leu Ser Leu Ser Asn Lys Phe Val Glu Gly
3395 3400 3405

Ser His Asn Ser Thr Val Ser Leu Thr Thr Lys Asn Met Glu Val
3410 3415 3420

Ser Val Ala Lys Thr Thr Lys Ala Glu Ile Pro Ile Leu Arg Met
3425 3430 3435

Asn Phe Lys Gln Glu Leu Asn Gly Asn Thr Lys Ser Lys Pro Thr

3440		3445		3450
Val Ser Ser Ser Met Glu Phe Lys Tyr Asp Phe Asn Ser Ser Met				
3455		3460		3465
Leu Tyr Ser Thr Ala Lys Gly Ala Val Asp His Lys Leu Ser Leu				
3470		3475		3480
Glu Ser Leu Thr Ser Tyr Phe Ser Ile Glu Ser Ser Thr Lys Gly				
3485		3490		3495
Asp Val Lys Gly Ser Val Leu Ser Arg Glu Tyr Ser Gly Thr Ile				
3500		3505		3510
Ala Ser Glu Ala Asn Thr Tyr Leu Asn Ser Lys Ser Thr Arg Ser				
3515		3520		3525
Ser Val Lys Leu Gln Gly Thr Ser Lys Ile Asp Asp Ile Trp Asn				
3530		3535		3540
Leu Glu Val Lys Glu Asn Phe Ala Gly Glu Ala Thr Leu Gln Arg				
3545		3550		3555
Ile Tyr Ser Leu Trp Glu His Ser Thr Lys Asn His Leu Gln Leu				
3560		3565		3570
Glu Gly Leu Phe Phe Thr Asn Gly Glu His Thr Ser Lys Ala Thr				
3575		3580		3585
Leu Glu Leu Ser Pro Trp Gln Met Ser Ala Leu Val Gln Val His				
3590		3595		3600
Ala Ser Gln Pro Ser Ser Phe His Asp Phe Pro Asp Leu Gly Gln				
3605		3610		3615
Glu Val Ala Leu Asn Ala Asn Thr Lys Asn Gln Lys Ile Arg Trp				
3620		3625		3630
Lys Asn Glu Val Arg Ile His Ser Gly Ser Phe Gln Ser Gln Val				
3635		3640		3645
Glu Leu Ser Asn Asp Gln Glu Lys Ala His Leu Asp Ile Ala Gly				
3650		3655		3660

Ser Leu Glu Gly His Leu Arg Phe Leu Lys Asn Ile Ile Leu Pro
3665 3670 3675

Val Tyr Asp Lys Ser Leu Trp Asp Phe Leu Lys Leu Asp Val Thr
3680 3685 3690

Thr Ser Ile Gly Arg Arg Gln His Leu Arg Val Ser Thr Ala Phe
3695 3700 3705

Val Tyr Thr Lys Asn Pro Asn Gly Tyr Ser Phe Ser Ile Pro Val
3710 3715 3720

Lys Val Leu Ala Asp Lys Phe Ile Thr Pro Gly Leu Lys Leu Asn
3725 3730 3735

Asp Leu Asn Ser Val Leu Val Met Pro Thr Phe His Val Pro Phe
3740 3745 3750

Thr Asp Leu Gln Val Pro Ser Cys Lys Leu Asp Phe Arg Glu Ile
3755 3760 3765

Gln Ile Tyr Lys Lys Leu Arg Thr Ser Ser Phe Ala Leu Asn Leu
3770 3775 3780

Pro Thr Leu Pro Glu Val Lys Phe Pro Glu Val Asp Val Leu Thr
3785 3790 3795

Lys Tyr Ser Gln Pro Glu Asp Ser Leu Ile Pro Phe Phe Glu Ile
3800 3805 3810

Thr Val Pro Glu Ser Gln Leu Thr Val Ser Gln Phe Thr Leu Pro
3815 3820 3825

Lys Ser Val Ser Asp Gly Ile Ala Ala Leu Asp Leu Asn Ala Val
3830 3835 3840

Ala Asn Lys Ile Ala Asp Phe Glu Leu Pro Thr Ile Ile Val Pro
3845 3850 3855

Glu Gln Thr Ile Glu Ile Pro Ser Ile Lys Phe Ser Val Pro Ala
3860 3865 3870

Gly Ile	Val Ile	Pro Ser	Phe	Gln Ala	Leu Thr	Ala	Arg Phe	Glu	
3875			3880			3885			
Val Asp	Ser Pro	Val Tyr	Asn	Ala Thr	Trp Ser	Ala	Ser Leu	Lys	
3890			3895			3900			
Asn Lys	Ala Asp	Tyr Val	Glu	Thr Val	Leu Asp	Ser	Thr Cys	Ser	
3905			3910			3915			
Ser Thr	Val Gln	Phe Leu	Glu	Tyr Glu	Leu Asn	Val	Leu Gly	Thr	
3920			3925			3930			
His Lys	Ile Glu	Asp Gly	Thr	Leu Ala	Ser Lys	Thr	Lys Gly	Thr	
3935			3940			3945			
Leu Ala	His Arg	Asp Phe	Ser	Ala Glu	Tyr Glu	Glu	Asp Gly	Lys	
3950			3955			3960			
Phe Glu	Gly Leu	Gln Glu	Trp	Glu Gly	Lys Ala	His	Leu Asn	Ile	
3965			3970			3975			
Lys Ser	Pro Ala	Phe Thr	Asp	Leu His	Leu Arg	Tyr	Gln Lys	Asp	
3980			3985			3990			
Lys Lys	Gly Ile	Ser Thr	Ser	Ala Ala	Ser Pro	Ala	Val Gly	Thr	
3995			4000			4005			
Val Gly	Met Asp	Met Asp	Glu	Asp Asp	Asp Phe	Ser	Lys Trp	Asn	
4010			4015			4020			
Phe Tyr	Tyr Ser	Pro Gln	Ser	Ser Pro	Asp Lys	Lys	Leu Thr	Ile	
4025			4030			4035			
Phe Lys	Thr Glu	Leu Arg	Val	Arg Glu	Ser Asp	Glu	Glu Thr	Gln	
4040			4045			4050			
Ile Lys	Val Asn	Trp Glu	Glu	Glu Ala	Ala Ser	Gly	Leu Leu	Thr	
4055			4060			4065			
Ser Leu	Lys Asp	Asn Val	Pro	Lys Ala	Thr Gly	Val	Leu Tyr	Asp	
4070			4075			4080			

Tyr Val Asn Lys Tyr His Trp Glu His Thr Gly Leu Thr Leu Arg
4085 4090 4095

Glu Val Ser Ser Lys Leu Arg Arg Asn Leu Gln Asn Asn Ala Glu
4100 4105 4110

Trp Val Tyr Gln Gly Ala Ile Arg Gln Ile Asp Asp Ile Asp Val
4115 4120 4125

Arg Phe Gln Lys Ala Ala Ser Gly Thr Thr Gly Thr Tyr Gln Glu
4130 4135 4140

Trp Lys Asp Lys Ala Gln Asn Leu Tyr Gln Glu Leu Leu Thr Gln
4145 4150 4155

Glu Gly Gln Ala Ser Phe Gln Gly Leu Lys Asp Asn Val Phe Asp
4160 4165 4170

Gly Leu Val Arg Val Thr Gln Lys Phe His Met Lys Val Lys His
4175 4180 4185

Leu Ile Asp Ser Leu Ile Asp Phe Leu Asn Phe Pro Arg Phe Gln
4190 4195 4200

Phe Pro Gly Lys Pro Gly Ile Tyr Thr Arg Glu Glu Leu Cys Thr
4205 4210 4215

Met Phe Ile Arg Glu Val Gly Thr Val Leu Ser Gln Val Tyr Ser
4220 4225 4230

Lys Val His Asn Gly Ser Glu Ile Leu Phe Ser Tyr Phe Gln Asp
4235 4240 4245

Leu Val Ile Thr Leu Pro Phe Glu Leu Arg Lys His Lys Leu Ile
4250 4255 4260

Asp Val Ile Ser Met Tyr Arg Glu Leu Leu Lys Asp Leu Ser Lys
4265 4270 4275

Glu Ala Gln Glu Val Phe Lys Ala Ile Gln Ser Leu Lys Thr Thr
4280 4285 4290

Glu Val Leu Arg Asn Leu Gln Asp Leu Leu Gln Phe Ile Phe Gln

4295		4300		4305
Leu Ile Glu Asp Asn Ile Lys Gln Leu Lys Glu Met Lys Phe Thr				
4310		4315		4320
Tyr Leu Ile Asn Tyr Ile Gln Asp Glu Ile Asn Thr Ile Phe Asn				
4325		4330		4335
Asp Tyr Ile Pro Tyr Val Phe Lys Leu Leu Lys Glu Asn Leu Cys				
4340		4345		4350
Leu Asn Leu His Lys Phe Asn Glu Phe Ile Gln Asn Glu Leu Gln				
4355		4360		4365
Glu Ala Ser Gln Glu Leu Gln Gln Ile His Gln Tyr Ile Met Ala				
4370		4375		4380
Leu Arg Glu Glu Tyr Phe Asp Pro Ser Ile Val Gly Trp Thr Val				
4385		4390		4395
Lys Tyr Tyr Glu Leu Glu Glu Lys Ile Val Ser Leu Ile Lys Asn				
4400		4405		4410
Leu Leu Val Ala Leu Lys Asp Phe His Ser Glu Tyr Ile Val Ser				
4415		4420		4425
Ala Ser Asn Phe Thr Ser Gln Leu Ser Ser Gln Val Glu Gln Phe				
4430		4435		4440
Leu His Arg Asn Ile Gln Glu Tyr Leu Ser Ile Leu Thr Asp Pro				
4445		4450		4455
Asp Gly Lys Gly Lys Glu Lys Ile Ala Glu Leu Ser Ala Thr Ala				
4460		4465		4470
Gln Glu Ile Ile Lys Ser Gln Ala Ile Ala Thr Lys Lys Ile Ile				
4475		4480		4485
Ser Asp Tyr His Gln Gln Phe Arg Tyr Lys Leu Gln Asp Phe Ser				
4490		4495		4500
Asp Gln Leu Ser Asp Tyr Tyr Glu Lys Phe Ile Ala Glu Ser Lys				
4505		4510		4515

Arg Leu Ile Asp Leu Ser Ile Gln Asn Tyr His Thr Phe Leu Ile
4520 4525 4530

Tyr Ile Thr Glu Leu Leu Lys Lys Leu Gln Ser Thr Thr Val Met
4535 4540 4545

Asn Pro Tyr Met Lys Leu Ala Pro Gly Glu Leu Thr Ile Ile Leu
4550 4555 4560

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<309> 1998-07-15
<313> (1)..(930)

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Met Lys Pro Pro Arg Pro Val Arg Thr Cys Ser Lys Val Leu Val Leu
1 5 10 15

Leu Ser Leu Leu Ala Ile His Gln Thr Thr Thr Ala Glu Lys Asn Gly
20 25 30

Ile Asp Ile Tyr Ser Leu Thr Val Asp Ser Arg Val Ser Ser Arg Phe
35 40 45

Ala His Thr Val Val Thr Ser Arg Val Val Asn Arg Ala Asn Thr Val
50 55 60

Gln Glu Ala Thr Phe Gln Met Glu Leu Pro Lys Lys Ala Phe Ile Thr
65 70 75 80

Asn Phe Ser Met Asn Ile Asp Gly Met Thr Tyr Pro Gly Ile Ile Lys
85 90 95

Glu Lys Ala Glu Ala Gln Ala Gln Tyr Ser Ala Ala Val Ala Lys Gly
100 105 110

Lys Ser Ala Gly Leu Val Lys Ala Thr Gly Arg Asn Met Glu Gln Phe
115 120 125

Gln Val Ser Val Ser Val Ala Pro Asn Ala Lys Ile Thr Phe Glu Leu
130 135 140

Val Tyr Glu Glu Leu Leu Lys Arg Arg Leu Gly Val Tyr Glu Leu Leu
145 150 155 160

Leu Lys Val Arg Pro Gln Gln Leu Val Lys His Leu Gln Met Asp Ile
165 170 175

His Ile Phe Glu Pro Gln Gly Ile Ser Phe Leu Glu Thr Glu Ser Thr
180 185 190

Phe Met Thr Asn Gln Leu Val Asp Ala Leu Thr Thr Trp Gln Asn Lys
195 200 205

Thr Lys Ala His Ile Arg Phe Lys Pro Thr Leu Ser Gln Gln Gln Lys
210 215 220

Ser Pro Glu Gln Gln Glu Thr Val Leu Asp Gly Asn Leu Ile Ile Arg
225 230 235 240

Tyr Asp Val Asp Arg Ala Ile Ser Gly Gly Ser Ile Gln Ile Glu Asn
245 250 255

Gly Tyr Phe Val His Tyr Phe Ala Pro Glu Gly Leu Thr Thr Met Pro
260 265 270

Lys Asn Val Val Phe Val Ile Asp Lys Ser Gly Ser Met Ser Gly Arg
275 280 285

Lys Ile Gln Gln Thr Arg Glu Ala Leu Ile Lys Ile Leu Asp Asp Leu
290 295 300

Ser Pro Arg Asp Gln Phe Asn Leu Ile Val Phe Ser Thr Glu Ala Thr
305 310 315 320

Gln Trp Arg Pro Ser Leu Val Pro Ala Ser Ala Glu Asn Val Asn Lys
325 330 335

Ala Arg Ser Phe Ala Ala Gly Ile Gln Ala Leu Gly Gly Thr Asn Ile
340 345 350

Asn Asp Ala Met Leu Met Ala Val Gln Leu Leu Asp Ser Ser Asn Gln
355 360 365

Glu Glu Arg Leu Pro Glu Gly Ser Val Ser Leu Ile Ile Leu Leu Thr
370 375 380

Asp Gly Asp Pro Thr Val Gly Glu Thr Asn Pro Arg Ser Ile Gln Asn
385 390 395 400

Asn Val Arg Glu Ala Val Ser Gly Arg Tyr Ser Leu Phe Cys Leu Gly
405 410 415

Phe Gly Phe Asp Val Ser Tyr Ala Phe Leu Glu Lys Leu Ala Leu Asp
420 425 430

Asn Gly Gly Leu Ala Arg Arg Ile His Glu Asp Ser Asp Ser Ala Leu
435 440 445

Gln Leu Gln Asp Phe Tyr Gln Glu Val Ala Asn Pro Leu Leu Thr Ala
450 455 460

Val Thr Phe Glu Tyr Pro Ser Asn Ala Val Glu Glu Val Thr Gln Asn
465 470 475 480

Asn Phe Arg Leu Leu Phe Lys Gly Ser Glu Met Val Val Ala Gly Lys
485 490 495

Leu Gln Asp Arg Gly Pro Asp Val Leu Thr Ala Thr Val Ser Gly Lys
500 505 510

Leu Pro Thr Gln Asn Ile Thr Phe Gln Thr Glu Ser Ser Val Ala Glu
515 520 525

Gln Glu Ala Glu Phe Gln Ser Pro Lys Tyr Ile Phe His Asn Phe Met
530 535 540

Glu Arg Leu Trp Ala Tyr Leu Thr Ile Gln Gln Leu Leu Glu Gln Thr
545 550 555 560

Val Ser Ala Ser Asp Ala Asp Gln Gln Ala Leu Arg Asn Gln Ala Leu
565 570 575

Asn Leu Ser Leu Ala Tyr Ser Phe Val Thr Pro Leu Thr Ser Met Val

580					585					590					
Val	Thr	Lys	Pro	Asp	Asp	Gln	Glu	Gln	Ser	Gln	Val	Ala	Glu	Lys	Pro
		595					600					605			
Met	Glu	Gly	Glu	Ser	Arg	Asn	Arg	Asn	Val	His	Ser	Gly	Ser	Thr	Phe
	610					615					620				
Phe	Lys	Tyr	Tyr	Leu	Gln	Gly	Ala	Lys	Ile	Pro	Lys	Pro	Glu	Ala	Ser
625					630					635					640
Phe	Ser	Pro	Arg	Arg	Gly	Trp	Asn	Arg	Gln	Ala	Gly	Ala	Ala	Gly	Ser
				645					650					655	
Arg	Met	Asn	Phe	Arg	Pro	Gly	Val	Leu	Ser	Ser	Arg	Gln	Leu	Gly	Leu
			660					665					670		
Pro	Gly	Pro	Pro	Asp	Val	Pro	Asp	His	Ala	Ala	Tyr	His	Pro	Phe	Arg
		675					680					685			
Arg	Leu	Ala	Ile	Leu	Pro	Ala	Ser	Ala	Pro	Pro	Ala	Thr	Ser	Asn	Pro
	690					695					700				
Asp	Pro	Ala	Val	Ser	Arg	Val	Met	Asn	Met	Lys	Ile	Glu	Glu	Thr	Thr
705					710					715					720
Met	Thr	Thr	Gln	Thr	Pro	Ala	Pro	Ile	Gln	Ala	Pro	Ser	Ala	Ile	Leu
			725						730					735	
Pro	Leu	Pro	Gly	Gln	Ser	Val	Glu	Arg	Leu	Cys	Val	Asp	Pro	Arg	His
			740					745					750		
Arg	Gln	Gly	Pro	Val	Asn	Leu	Leu	Ser	Asp	Pro	Glu	Gln	Gly	Val	Glu
		755					760					765			
Val	Thr	Gly	Gln	Tyr	Glu	Arg	Glu	Lys	Ala	Gly	Phe	Ser	Trp	Ile	Glu
	770					775					780				
Val	Thr	Phe	Lys	Asn	Pro	Leu	Val	Trp	Val	His	Ala	Ser	Pro	Glu	His
785					790					795					800
Val	Val	Val	Thr	Arg	Asn	Arg	Arg	Ser	Ser	Ala	Tyr	Lys	Trp	Lys	Glu
				805					810					815	

Thr Leu Phe Ser Val Met Pro Gly Leu Lys Met Thr Met Asp Lys Thr
820 825 830

Gly Leu Leu Leu Leu Ser Asp Pro Asp Lys Val Thr Ile Gly Leu Leu
835 840 845

Phe Trp Asp Gly Arg Gly Glu Gly Leu Arg Leu Leu Leu Arg Asp Thr
850 855 860

Asp Arg Phe Ser Ser His Val Gly Gly Thr Leu Gly Gln Phe Tyr Gln
865 870 875 880

Glu Val Leu Trp Gly Ser Pro Ala Ala Ser Asp Asp Gly Arg Arg Thr
885 890 895

Leu Arg Val Gln Gly Asn Asp His Ser Ala Thr Arg Glu Arg Arg Leu
900 905 910

Asp Tyr Gln Glu Gly Pro Pro Gly Val Glu Ile Ser Cys Trp Ser Val
915 920 925

Glu Leu
930

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<309> 1986-07-21
<313> (1)..(1744)

<400> 43

Met Arg Leu Leu Trp Gly Leu Ile Trp Ala Ser Ser Phe Phe Thr Leu
1 5 10 15

Ser Leu Gln Lys Pro Arg Leu Leu Leu Phe Ser Pro Ser Val Val His
20 25 30

Leu Gly Val Pro Leu Ser Val Gly Val Gln Leu Gln Asp Val Pro Arg
35 40 45

Gly Gln Val Val Lys Gly Ser Val Phe Leu Arg Asn Pro Ser Arg Asn
50 55 60

Asn Val Pro Cys Ser Pro Lys Val Asp Phe Thr Leu Ser Ser Glu Arg
65 70 75 80

Asp Phe Ala Leu Leu Ser Leu Gln Val Pro Leu Lys Asp Ala Lys Ser
85 90 95

Cys Gly Leu His Gln Leu Leu Arg Gly Pro Glu Val Gln Leu Val Ala
100 105 110

His Ser Pro Trp Leu Lys Asp Ser Leu Ser Arg Thr Thr Asn Ile Gln
115 120 125

Gly Ile Asn Leu Leu Phe Ser Ser Arg Arg Gly His Leu Phe Leu Gln
130 135 140

Thr Asp Gln Pro Ile Tyr Asn Pro Gly Gln Arg Val Arg Tyr Arg Val
145 150 155 160

Phe Ala Leu Asp Gln Lys Met Arg Pro Ser Thr Asp Thr Ile Thr Val
165 170 175

Met Val Glu Asn Ser His Gly Leu Arg Val Arg Lys Lys Glu Val Tyr
180 185 190

Met Pro Ser Ser Ile Phe Gln Asp Asp Phe Val Ile Pro Asp Ile Ser
195 200 205

Glu Pro Gly Thr Trp Lys Ile Ser Ala Arg Phe Ser Asp Gly Leu Glu
210 215 220

Ser Asn Ser Ser Thr Gln Phe Glu Val Lys Lys Tyr Val Leu Pro Asn
225 230 235 240

Phe Glu Val Lys Ile Thr Pro Gly Lys Pro Tyr Ile Leu Thr Val Pro
245 250 255

Gly His Leu Asp Glu Met Gln Leu Asp Ile Gln Ala Arg Tyr Ile Tyr
260 265 270

Gly Lys Pro Val Gln Gly Val Ala Tyr Val Arg Phe Gly Leu Leu Asp
275 280 285

Glu Asp Gly Lys Lys Thr Phe Phe Arg Gly Leu Glu Ser Gln Thr Lys
290 295 300

Leu Val Asn Gly Gln Ser His Ile Ser Leu Ser Lys Ala Glu Phe Gln
305 310 315 320

Asp Ala Leu Glu Lys Leu Asn Met Gly Ile Thr Asp Leu Gln Gly Leu
325 330 335

Arg Leu Tyr Val Ala Ala Ala Ile Ile Glu Ser Pro Gly Gly Glu Met
340 345 350

Glu Glu Ala Glu Leu Thr Ser Trp Tyr Phe Val Ser Ser Pro Phe Ser
355 360 365

Leu Asp Leu Ser Lys Thr Lys Arg His Leu Val Pro Gly Ala Pro Phe
370 375 380

Leu Leu Gln Ala Leu Val Arg Glu Met Ser Gly Ser Pro Ala Ser Gly
385 390 395 400

Ile Pro Val Lys Val Ser Ala Thr Val Ser Ser Pro Gly Ser Val Pro
405 410 415

Glu Val Gln Asp Ile Gln Gln Asn Thr Asp Gly Ser Gly Gln Val Ser
420 425 430

Ile Pro Ile Ile Ile Pro Gln Thr Ile Ser Glu Leu Gln Leu Ser Val
435 440 445

Ser Ala Gly Ser Pro His Pro Ala Ile Ala Arg Leu Thr Val Ala Ala
450 455 460

Pro Pro Ser Gly Gly Pro Gly Phe Leu Ser Ile Glu Arg Pro Asp Ser
465 470 475 480

Arg Pro Pro Arg Val Gly Asp Thr Leu Asn Leu Asn Leu Arg Ala Val
485 490 495

Gly Ser Gly Ala Thr Phe Ser His Tyr Tyr Tyr Met Ile Leu Ser Arg

500					505					510					
Gly	Gln	Ile	Val	Phe	Met	Asn	Arg	Glu	Pro	Lys	Arg	Thr	Leu	Thr	Ser
		515					520					525			
Val	Ser	Val	Phe	Val	Asp	His	His	Leu	Ala	Pro	Ser	Phe	Tyr	Phe	Val
	530					535					540				
Ala	Phe	Tyr	Tyr	His	Gly	Asp	His	Pro	Val	Ala	Asn	Ser	Leu	Arg	Val
545					550					555					560
Asp	Val	Gln	Ala	Gly	Ala	Cys	Glu	Gly	Lys	Leu	Glu	Leu	Ser	Val	Asp
				565					570					575	
Gly	Ala	Lys	Gln	Tyr	Arg	Asn	Gly	Glu	Ser	Val	Lys	Leu	His	Leu	Glu
			580					585					590		
Thr	Asp	Ser	Leu	Ala	Leu	Val	Ala	Leu	Gly	Ala	Leu	Asp	Thr	Ala	Leu
		595					600					605			
Tyr	Ala	Ala	Gly	Ser	Lys	Ser	His	Lys	Pro	Leu	Asn	Met	Gly	Lys	Val
	610					615					620				
Phe	Glu	Ala	Met	Asn	Ser	Tyr	Asp	Leu	Gly	Cys	Gly	Pro	Gly	Gly	Gly
625					630					635					640
Asp	Ser	Ala	Leu	Gln	Val	Phe	Gln	Ala	Ala	Gly	Leu	Ala	Phe	Ser	Asp
				645					650					655	
Gly	Asp	Gln	Trp	Thr	Leu	Ser	Arg	Lys	Arg	Leu	Ser	Cys	Pro	Lys	Glu
		660						665					670		
Lys	Thr	Thr	Arg	Lys	Lys	Arg	Asn	Val	Asn	Phe	Gln	Lys	Ala	Ile	Asn
		675					680					685			
Glu	Lys	Leu	Gly	Gln	Tyr	Ala	Ser	Pro	Thr	Ala	Lys	Arg	Cys	Cys	Gln
	690					695					700				
Asp	Gly	Val	Thr	Arg	Leu	Pro	Met	Met	Arg	Ser	Cys	Glu	Gln	Arg	Ala
705					710					715					720
Ala	Arg	Val	Gln	Gln	Pro	Asp	Cys	Arg	Glu	Pro	Phe	Leu	Ser	Cys	Cys
			725						730					735	

Gln Phe Ala Glu Ser Leu Arg Lys Lys Ser Arg Asp Lys Gly Gln Ala
740 745 750

Gly Leu Gln Arg Ala Leu Glu Ile Leu Gln Glu Glu Asp Leu Ile Asp
755 760 765

Glu Asp Asp Ile Pro Val Arg Ser Phe Phe Pro Glu Asn Trp Leu Trp
770 775 780

Arg Val Glu Thr Val Asp Arg Phe Gln Ile Leu Thr Leu Trp Leu Pro
785 790 795 800

Asp Ser Leu Thr Thr Trp Glu Ile His Gly Leu Ser Leu Ser Lys Thr
805 810 815

Lys Gly Leu Cys Val Ala Thr Pro Val Gln Leu Arg Val Phe Arg Glu
820 825 830

Phe His Leu His Leu Arg Leu Pro Met Ser Val Arg Arg Phe Glu Gln
835 840 845

Leu Glu Leu Arg Pro Val Leu Tyr Asn Tyr Leu Asp Lys Asn Leu Thr
850 855 860

Val Ser Val His Val Ser Pro Val Glu Gly Leu Cys Leu Ala Gly Gly
865 870 875 880

Gly Gly Leu Ala Gln Gln Val Leu Val Pro Ala Gly Ser Ala Arg Pro
885 890 895

Val Ala Phe Ser Val Val Pro Thr Ala Ala Ala Val Ser Leu Lys
900 905 910

Val Val Ala Arg Gly Ser Phe Glu Phe Pro Val Gly Asp Ala Val Ser
915 920 925

Lys Val Leu Gln Ile Glu Lys Glu Gly Ala Ile His Arg Glu Glu Leu
930 935 940

Val Tyr Glu Leu Asn Pro Leu Asp His Arg Gly Arg Thr Leu Glu Ile
945 950 955 960

Pro Gly Asn Ser Asp Pro Asn Met Ile Pro Asp Gly Asp Phe Asn Ser
965 970 975

Tyr Val Arg Val Thr Ala Ser Asp Pro Leu Asp Thr Leu Gly Ser Glu
980 985 990

Gly Ala Leu Ser Pro Gly Gly Val Ala Ser Leu Leu Arg Leu Pro Arg
995 1000 1005

Gly Cys Gly Glu Gln Thr Met Ile Tyr Leu Ala Pro Thr Leu Ala
1010 1015 1020

Ala Ser Arg Tyr Leu Asp Lys Thr Glu Gln Trp Ser Thr Leu Pro
1025 1030 1035

Pro Glu Thr Lys Asp His Ala Val Asp Leu Ile Gln Lys Gly Tyr
1040 1045 1050

Met Arg Ile Gln Gln Phe Arg Lys Ala Asp Gly Ser Tyr Ala Ala
1055 1060 1065

Trp Leu Ser Arg Asp Ser Ser Thr Trp Leu Thr Ala Phe Val Leu
1070 1075 1080

Lys Val Leu Ser Leu Ala Gln Glu Gln Val Gly Gly Ser Pro Glu
1085 1090 1095

Lys Leu Gln Glu Thr Ser Asn Trp Leu Leu Ser Gln Gln Gln Ala
1100 1105 1110

Asp Gly Ser Phe Gln Asp Pro Cys Pro Val Leu Asp Arg Ser Met
1115 1120 1125

Gln Gly Gly Leu Val Gly Asn Asp Glu Thr Val Ala Leu Thr Ala
1130 1135 1140

Phe Val Thr Ile Ala Leu His His Gly Leu Ala Val Phe Gln Asp
1145 1150 1155

Glu Gly Ala Glu Pro Leu Lys Gln Arg Val Glu Ala Ser Ile Ser
1160 1165 1170

Lys	Ala	Asn	Ser	Phe	Leu	Gly	Glu	Lys	Ala	Ser	Ala	Gly	Leu	Leu
1175						1180					1185			
Gly	Ala	His	Ala	Ala	Ala	Ile	Thr	Ala	Tyr	Ala	Leu	Ser	Leu	Thr
1190						1195					1200			
Lys	Ala	Pro	Val	Asp	Leu	Leu	Gly	Val	Ala	His	Asn	Asn	Leu	Met
1205						1210					1215			
Ala	Met	Ala	Gln	Glu	Thr	Gly	Asp	Asn	Leu	Tyr	Trp	Gly	Ser	Val
1220						1225					1230			
Thr	Gly	Ser	Gln	Ser	Asn	Ala	Val	Ser	Pro	Thr	Pro	Ala	Pro	Arg
1235						1240					1245			
Asn	Pro	Ser	Asp	Pro	Met	Pro	Gln	Ala	Pro	Ala	Leu	Trp	Ile	Glu
1250						1255					1260			
Thr	Thr	Ala	Tyr	Ala	Leu	Leu	His	Leu	Leu	Leu	His	Glu	Gly	Lys
1265						1270					1275			
Ala	Glu	Met	Ala	Asp	Gln	Ala	Ser	Ala	Trp	Leu	Thr	Arg	Gln	Gly
1280						1285					1290			
Ser	Phe	Gln	Gly	Gly	Phe	Arg	Ser	Thr	Gln	Asp	Thr	Val	Ile	Ala
1295						1300					1305			
Leu	Asp	Ala	Leu	Ser	Ala	Tyr	Trp	Ile	Ala	Ser	His	Thr	Thr	Glu
1310						1315					1320			
Glu	Arg	Gly	Leu	Asn	Val	Thr	Leu	Ser	Ser	Thr	Gly	Arg	Asn	Gly
1325						1330					1335			
Phe	Lys	Ser	His	Ala	Leu	Gln	Leu	Asn	Asn	Arg	Gln	Ile	Arg	Gly
1340						1345					1350			
Leu	Glu	Glu	Glu	Leu	Gln	Phe	Ser	Leu	Gly	Ser	Lys	Ile	Asn	Val
1355						1360					1365			
Lys	Val	Gly	Gly	Asn	Ser	Lys	Gly	Thr	Leu	Lys	Val	Leu	Arg	Thr
1370						1375					1380			
Tyr	Asn	Val	Leu	Asp	Met	Lys	Asn	Thr	Thr	Cys	Gln	Asp	Leu	Gln

1385		1390		1395
Ile Glu Val Thr Val Lys Gly His Val Glu Tyr Thr Met Glu Ala				
1400		1405		1410
Asn Glu Asp Tyr Glu Asp Tyr Glu Tyr Asp Glu Leu Pro Ala Lys				
1415		1420		1425
Asp Asp Pro Asp Ala Pro Leu Gln Pro Val Thr Pro Leu Gln Leu				
1430		1435		1440
Phe Glu Gly Arg Arg Asn Arg Arg Arg Arg Glu Ala Pro Lys Val				
1445		1450		1455
Val Glu Glu Gln Glu Ser Arg Val His Tyr Thr Val Cys Ile Trp				
1460		1465		1470
Arg Asn Gly Lys Val Gly Leu Ser Gly Met Ala Ile Ala Asp Val				
1475		1480		1485
Thr Leu Leu Ser Gly Phe His Ala Leu Arg Ala Asp Leu Glu Lys				
1490		1495		1500
Leu Thr Ser Leu Ser Asp Arg Tyr Val Ser His Phe Glu Thr Glu				
1505		1510		1515
Gly Pro His Val Leu Leu Tyr Phe Asp Ser Val Pro Thr Ser Arg				
1520		1525		1530
Glu Cys Val Gly Phe Glu Ala Val Gln Glu Val Pro Val Gly Leu				
1535		1540		1545
Val Gln Pro Ala Ser Ala Thr Leu Tyr Asp Tyr Tyr Asn Pro Glu				
1550		1555		1560
Arg Arg Cys Ser Val Phe Tyr Gly Ala Pro Ser Lys Ser Arg Leu				
1565		1570		1575
Leu Ala Thr Leu Cys Ser Ala Glu Val Cys Gln Cys Ala Glu Gly				
1580		1585		1590
Lys Cys Pro Arg Gln Arg Arg Ala Leu Glu Arg Gly Leu Gln Asp				
1595		1600		1605

Glu Asp Gly Tyr Arg Met Lys Phe Ala Cys Tyr Tyr Pro Arg Val
 1610 1615 1620

Glu Tyr Gly Phe Gln Val Lys Val Leu Arg Glu Asp Ser Arg Ala
 1625 1630 1635

Ala Phe Arg Leu Phe Glu Thr Lys Ile Thr Gln Val Leu His Phe
 1640 1645 1650

Thr Lys Asp Val Lys Ala Ala Ala Asn Gln Met Arg Asn Phe Leu
 1655 1660 1665

Val Arg Ala Ser Cys Arg Leu Arg Leu Glu Pro Gly Lys Glu Tyr
 1670 1675 1680

Leu Ile Met Gly Leu Asp Gly Ala Thr Tyr Asp Leu Glu Gly His
 1685 1690 1695

Pro Gln Tyr Leu Leu Asp Ser Asn Ser Trp Ile Glu Glu Met Pro
 1700 1705 1710

Ser Glu Arg Leu Cys Arg Ser Thr Arg Gln Arg Ala Ala Cys Ala
 1715 1720 1725

Gln Leu Asn Asp Phe Leu Gln Glu Tyr Gly Thr Gln Gly Cys Gln
 1730 1735 1740

Val

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 <212> PRT
 <213> Homo sapiens

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 <309> 1986-07-21
 <313> (1)..(1663)

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Leu Pro Leu Ala Leu Gly Ser Pro Met Tyr Ser Ile Ile Thr Pro Asn
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Ile Leu Arg Leu Glu Ser Glu Glu Thr Met Val Leu Glu Ala His Asp
35 40 45

Ala Gln Gly Asp Val Pro Val Thr Val Thr Val His Asp Phe Pro Gly
50 55 60

Lys Lys Leu Val Leu Ser Ser Glu Lys Thr Val Leu Thr Pro Ala Thr
65 70 75 80

Asn His Met Gly Asn Val Thr Phe Thr Ile Pro Ala Asn Arg Glu Phe
85 90 95

Lys Ser Glu Lys Gly Arg Asn Lys Phe Val Thr Val Gln Ala Thr Phe
100 105 110

Gly Thr Gln Val Val Glu Lys Val Val Leu Val Ser Leu Gln Ser Gly
115 120 125

Tyr Leu Phe Ile Gln Thr Asp Lys Thr Ile Tyr Thr Pro Gly Ser Thr
130 135 140

Val Leu Tyr Arg Ile Phe Thr Val Asn His Lys Leu Leu Pro Val Gly
145 150 155 160

Arg Thr Val Met Val Asn Ile Glu Asn Pro Glu Gly Ile Pro Val Lys
165 170 175

Gln Asp Ser Leu Ser Ser Gln Asn Gln Leu Gly Val Leu Pro Leu Ser
180 185 190

Trp Asp Ile Pro Glu Leu Val Asn Met Gly Gln Trp Lys Ile Arg Ala
195 200 205

Tyr Tyr Glu Asn Ser Pro Gln Gln Val Phe Ser Thr Glu Phe Glu Val
210 215 220

Lys Glu Tyr Val Leu Pro Ser Phe Glu Val Ile Val Glu Pro Thr Glu
225 230 235 240

Lys Phe Tyr Tyr Ile Tyr Asn Glu Lys Gly Leu Glu Val Thr Ile Thr
245 250 255

Ala Arg Phe Leu Tyr Gly Lys Lys Val Glu Gly Thr Ala Phe Val Ile
260 265 270

Phe Gly Ile Gln Asp Gly Glu Gln Arg Ile Ser Leu Pro Glu Ser Leu
275 280 285

Lys Arg Ile Pro Ile Glu Asp Gly Ser Gly Glu Val Val Leu Ser Arg
290 295 300

Lys Val Leu Leu Asp Gly Val Gln Asn Leu Arg Ala Glu Asp Leu Val
305 310 315 320

Gly Lys Ser Leu Tyr Val Ser Ala Thr Val Ile Leu His Ser Gly Ser
325 330 335

Asp Met Val Gln Ala Glu Arg Ser Gly Ile Pro Ile Val Thr Ser Pro
340 345 350

Tyr Gln Ile His Phe Thr Lys Thr Pro Lys Tyr Phe Lys Pro Gly Met
355 360 365

Pro Phe Asp Leu Met Val Phe Val Thr Asn Pro Asp Gly Ser Pro Ala
370 375 380

Tyr Arg Val Pro Val Ala Val Gln Gly Glu Asp Thr Val Gln Ser Leu
385 390 395 400

Thr Gln Gly Asp Gly Val Ala Lys Leu Ser Ile Asn Thr His Pro Ser
405 410 415

Gln Lys Pro Leu Ser Ile Thr Val Arg Thr Lys Lys Gln Glu Leu Ser
420 425 430

Glu Ala Glu Gln Ala Thr Arg Thr Met Gln Ala Leu Pro Tyr Ser Thr
435 440 445

Val Gly Asn Ser Asn Asn Tyr Leu His Leu Ser Val Leu Arg Thr Glu
450 455 460

Leu Arg Pro Gly Glu Thr Leu Asn Val Asn Phe Leu Leu Arg Met Asp

465		470		475		480									
Arg	Ala	His	Glu	Ala	Lys	Ile	Arg	Tyr	Tyr	Thr	Tyr	Leu	Ile	Met	Asn
			485					490						495	
Lys	Gly	Arg	Leu	Leu	Lys	Ala	Gly	Arg	Gln	Val	Arg	Glu	Pro	Gly	Gln
			500					505					510		
Asp	Leu	Val	Val	Leu	Pro	Leu	Ser	Ile	Thr	Thr	Asp	Phe	Ile	Pro	Ser
		515					520					525			
Phe	Arg	Leu	Val	Ala	Tyr	Tyr	Thr	Leu	Ile	Gly	Ala	Ser	Gly	Gln	Arg
	530					535					540				
Glu	Val	Val	Ala	Asp	Ser	Val	Trp	Val	Asp	Val	Lys	Asp	Ser	Cys	Val
545					550				555					560	
Gly	Ser	Leu	Val	Val	Lys	Ser	Gly	Gln	Ser	Glu	Asp	Arg	Gln	Pro	Val
			565					570						575	
Pro	Gly	Gln	Gln	Met	Thr	Leu	Lys	Ile	Glu	Gly	Asp	His	Gly	Ala	Arg
			580					585					590		
Val	Val	Leu	Val	Ala	Val	Asp	Lys	Gly	Val	Phe	Val	Leu	Asn	Lys	Lys
		595					600					605			
Asn	Lys	Leu	Thr	Gln	Ser	Lys	Ile	Trp	Asp	Val	Val	Glu	Lys	Ala	Asp
	610					615						620			
Ile	Gly	Cys	Thr	Pro	Gly	Ser	Gly	Lys	Asp	Tyr	Ala	Gly	Val	Phe	Ser
625					630				635					640	
Asp	Ala	Gly	Leu	Thr	Phe	Thr	Ser	Ser	Ser	Gly	Gln	Gln	Thr	Ala	Gln
				645					650					655	
Arg	Ala	Glu	Leu	Gln	Cys	Pro	Gln	Pro	Ala	Ala	Arg	Arg	Arg	Arg	Ser
			660						665					670	
Val	Gln	Leu	Thr	Glu	Lys	Arg	Met	Asp	Lys	Val	Gly	Lys	Tyr	Pro	Lys
		675					680					685			
Glu	Leu	Arg	Lys	Cys	Cys	Glu	Asp	Gly	Met	Arg	Glu	Asn	Pro	Met	Arg
	690					695					700				

Phe Ser Cys Gln Arg Arg Thr Arg Phe Ile Ser Leu Gly Glu Ala Cys
705 710 715 720

Lys Lys Val Phe Leu Asp Cys Cys Asn Tyr Ile Thr Glu Leu Arg Arg
725 730 735

Gln His Ala Arg Ala Ser His Leu Gly Leu Ala Arg Ser Asn Leu Asp
740 745 750

Glu Asp Ile Ile Ala Glu Glu Asn Ile Val Ser Arg Ser Glu Phe Pro
755 760 765

Glu Ser Trp Leu Trp Asn Val Glu Asp Leu Lys Glu Pro Pro Lys Asn
770 775 780

Gly Ile Ser Thr Lys Leu Met Asn Ile Phe Leu Lys Asp Ser Ile Thr
785 790 795 800

Thr Trp Glu Ile Leu Ala Val Ser Met Ser Asp Lys Lys Gly Ile Cys
805 810 815

Val Ala Asp Pro Phe Glu Val Thr Val Met Gln Asp Phe Phe Ile Asp
820 825 830

Leu Arg Leu Pro Tyr Ser Val Val Arg Asn Glu Gln Val Glu Ile Arg
835 840 845

Ala Val Leu Tyr Asn Tyr Arg Gln Asn Gln Glu Leu Lys Val Arg Val
850 855 860

Glu Leu Leu His Asn Pro Ala Phe Cys Ser Leu Ala Thr Thr Lys Arg
865 870 875 880

Arg His Gln Gln Thr Val Thr Ile Pro Pro Lys Ser Ser Leu Ser Val
885 890 895

Pro Tyr Val Ile Val Pro Leu Lys Thr Gly Leu Gln Glu Val Glu Val
900 905 910

Lys Ala Ala Val Tyr His His Phe Ile Ser Asp Gly Val Arg Lys Ser
915 920 925

Leu Lys Val Val Pro Glu Gly Ile Arg Met Asn Lys Thr Val Ala Val
930 935 940

Arg Thr Leu Asp Pro Glu Arg Leu Gly Arg Glu Gly Val Gln Lys Glu
945 950 955 960

Asp Ile Pro Pro Ala Asp Leu Ser Asp Gln Val Pro Asp Thr Glu Ser
965 970 975

Glu Thr Arg Ile Leu Leu Gln Gly Thr Pro Val Ala Gln Met Thr Glu
980 985 990

Asp Ala Val Asp Ala Glu Arg Leu Lys His Leu Ile Val Thr Pro Ser
995 1000 1005

Gly Cys Gly Glu Gln Asn Met Ile Gly Met Thr Pro Thr Val Ile
1010 1015 1020

Ala Val His Tyr Leu Asp Glu Thr Glu Gln Trp Glu Lys Phe Gly
1025 1030 1035

Leu Glu Lys Arg Gln Gly Ala Leu Glu Leu Ile Lys Lys Gly Tyr
1040 1045 1050

Thr Gln Gln Leu Ala Phe Arg Gln Pro Ser Ser Ala Phe Ala Ala
1055 1060 1065

Phe Val Lys Arg Ala Pro Ser Thr Trp Leu Thr Ala Tyr Val Val
1070 1075 1080

Lys Val Phe Ser Leu Ala Val Asn Leu Ile Ala Ile Asp Ser Gln
1085 1090 1095

Val Leu Cys Gly Ala Val Lys Trp Leu Ile Leu Glu Lys Gln Lys
1100 1105 1110

Pro Asp Gly Val Phe Gln Glu Asp Ala Pro Val Ile His Gln Glu
1115 1120 1125

Met Ile Gly Gly Leu Arg Asn Asn Asn Glu Lys Asp Met Ala Leu
1130 1135 1140

Thr	Ala	Phe	Val	Leu	Ile	Ser	Leu	Gln	Glu	Ala	Lys	Asp	Ile	Cys
1145						1150					1155			
Glu	Glu	Gln	Val	Asn	Ser	Leu	Pro	Gly	Ser	Ile	Thr	Lys	Ala	Gly
1160						1165					1170			
Asp	Phe	Leu	Glu	Ala	Asn	Tyr	Met	Asn	Leu	Gln	Arg	Ser	Tyr	Thr
1175						1180					1185			
Val	Ala	Ile	Ala	Gly	Tyr	Ala	Leu	Ala	Gln	Met	Gly	Arg	Leu	Lys
1190						1195					1200			
Gly	Pro	Leu	Leu	Asn	Lys	Phe	Leu	Thr	Thr	Ala	Lys	Asp	Lys	Asn
1205						1210					1215			
Arg	Trp	Glu	Asp	Pro	Gly	Lys	Gln	Leu	Tyr	Asn	Val	Glu	Ala	Thr
1220						1225					1230			
Ser	Tyr	Ala	Leu	Leu	Ala	Leu	Leu	Gln	Leu	Lys	Asp	Phe	Asp	Phe
1235						1240					1245			
Val	Pro	Pro	Val	Val	Arg	Trp	Leu	Asn	Glu	Gln	Arg	Tyr	Tyr	Gly
1250						1255					1260			
Gly	Gly	Tyr	Gly	Ser	Thr	Gln	Ala	Thr	Phe	Met	Val	Phe	Gln	Ala
1265						1270					1275			
Leu	Ala	Gln	Tyr	Gln	Lys	Asp	Ala	Pro	Asp	His	Gln	Glu	Leu	Asn
1280						1285					1290			
Leu	Asp	Val	Ser	Leu	Gln	Leu	Pro	Ser	Arg	Ser	Ser	Lys	Ile	Thr
1295						1300					1305			
His	Arg	Ile	His	Trp	Glu	Ser	Ala	Ser	Leu	Leu	Arg	Ser	Glu	Glu
1310						1315					1320			
Thr	Lys	Glu	Asn	Glu	Gly	Phe	Thr	Val	Thr	Ala	Glu	Gly	Lys	Gly
1325						1330					1335			
Gln	Gly	Thr	Leu	Ser	Val	Val	Thr	Met	Tyr	His	Ala	Lys	Ala	Lys
1340						1345					1350			
Asp	Gln	Leu	Thr	Cys	Asn	Lys	Phe	Asp	Leu	Lys	Val	Thr	Ile	Lys

1355		1360		1365
Pro Ala	Pro Glu Thr Glu Lys	Arg Pro Gln Asp	Ala Lys Asn Thr	
1370	1375		1380	
Met Ile	Leu Glu Ile Cys Thr	Arg Tyr Arg Gly Asp	Gln Asp Ala	
1385	1390		1395	
Thr Met	Ser Ile Leu Asp Ile	Ser Met Met Thr Gly	Phe Ala Pro	
1400	1405		1410	
Asp Thr	Asp Asp Leu Lys Gln	Leu Ala Asn Gly Val	Asp Arg Tyr	
1415	1420		1425	
Ile Ser	Lys Tyr Glu Leu Asp	Lys Ala Phe Ser Asp	Arg Asn Thr	
1430	1435		1440	
Leu Ile	Ile Tyr Leu Asp Lys	Val Ser His Ser Glu	Asp Asp Cys	
1445	1450		1455	
Leu Ala	Phe Lys Val His Gln	Tyr Phe Asn Val Glu	Leu Ile Gln	
1460	1465		1470	
Pro Gly	Ala Val Lys Val Tyr	Ala Tyr Tyr Asn Leu	Glu Glu Ser	
1475	1480		1485	
Cys Thr	Arg Phe Tyr His Pro	Glu Lys Glu Asp Gly	Lys Leu Asn	
1490	1495		1500	
Lys Leu	Cys Arg Asp Glu Leu	Cys Arg Cys Ala Glu	Glu Asn Cys	
1505	1510		1515	
Phe Ile	Gln Lys Ser Asp Asp	Lys Val Thr Leu Glu	Glu Arg Leu	
1520	1525		1530	
Asp Lys	Ala Cys Glu Pro Gly	Val Asp Tyr Val Tyr	Lys Thr Arg	
1535	1540		1545	
Leu Val	Lys Val Gln Leu Ser	Asn Asp Phe Asp Glu	Tyr Ile Met	
1550	1555		1560	
Ala Ile	Glu Gln Thr Ile Lys	Ser Gly Ser Asp Glu	Val Gln Val	
1565	1570		1575	

Gly Gln Gln Arg Thr Phe Ile Ser Pro Ile Lys Cys Arg Glu Ala
1580 1585 1590

Leu Lys Leu Glu Glu Lys Lys His Tyr Leu Met Trp Gly Leu Ser
1595 1600 1605

Ser Asp Phe Trp Gly Glu Lys Pro Asn Leu Ser Tyr Ile Ile Gly
1610 1615 1620

Lys Asp Thr Trp Val Glu His Trp Pro Glu Glu Asp Glu Cys Gln
1625 1630 1635

Asp Glu Glu Asn Gln Lys Gln Cys Gln Asp Leu Gly Ala Phe Thr
1640 1645 1650

Glu Ser Met Val Val Phe Gly Cys Pro Asn
1655 1660

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<300>
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<309> 2003-02-28
<313> (1)..(93)

<400> 45

Met Ser Gly Leu Arg Val Tyr Ser Thr Ser Val Thr Gly Ser Arg Glu
1 5 10 15

Ile Lys Ser Gln Gln Ser Glu Val Thr Arg Ile Leu Asp Gly Lys Arg
20 25 30

Ile Gln Tyr Gln Leu Val Asp Ile Ser Gln Asp Asn Ala Leu Arg Asp
35 40 45

Glu Met Arg Ala Leu Ala Gly Asn Pro Lys Ala Thr Pro Pro Gln Ile
50 55 60

Val Asn Gly Asp Gln Tyr Cys Gly Asp Tyr Glu Leu Phe Val Glu Ala
65 70 75 80

Val Glu Gln Asn Thr Leu Gln Glu Phe Leu Lys Leu Ala
85 90

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<211> 567
<212> PRT
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<300>
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<309> 2003-02-28
<313> (1)..(567)

<400> 46

Met Ala Pro Leu Ala Leu His Leu Leu Val Leu Val Pro Ile Leu Leu
1 5 10 15

Ser Leu Val Ala Ser Gln Asp Trp Lys Ala Glu Arg Ser Gln Asp Pro
20 25 30

Phe Glu Lys Cys Met Gln Asp Pro Asp Tyr Glu Gln Leu Leu Lys Val
35 40 45

Val Thr Trp Gly Leu Asn Arg Thr Leu Lys Pro Gln Arg Val Ile Val
50 55 60

Val Gly Ala Gly Val Ala Gly Leu Val Ala Ala Lys Val Leu Ser Asp
65 70 75 80

Ala Gly His Lys Val Thr Ile Leu Glu Ala Asp Asn Arg Ile Gly Gly
85 90 95

Arg Ile Phe Thr Tyr Arg Asp Gln Asn Thr Gly Trp Ile Gly Glu Leu
100 105 110

Gly Ala Met Arg Met Pro Ser Ser His Arg Ile Leu His Lys Leu Cys
115 120 125

Gln Gly Leu Gly Leu Asn Leu Thr Lys Phe Thr Gln Tyr Asp Lys Asn
130 135 140

Thr Trp Thr Glu Val His Glu Val Lys Leu Arg Asn Tyr Val Val Glu
145 150 155 160

Lys Val Pro Glu Lys Leu Gly Tyr Ala Leu Arg Pro Gln Glu Lys Gly
165 170 175

His Ser Pro Glu Asp Ile Tyr Gln Met Ala Leu Asn Gln Ala Leu Lys
180 185 190

Asp Leu Lys Ala Leu Gly Cys Arg Lys Ala Met Lys Lys Phe Glu Arg
195 200 205

His Thr Leu Leu Glu Tyr Leu Leu Gly Glu Gly Asn Leu Ser Arg Pro
210 215 220

Ala Val Gln Leu Leu Gly Asp Val Met Ser Glu Asp Gly Phe Phe Tyr
225 230 235 240

Leu Ser Phe Ala Glu Ala Leu Arg Ala His Ser Cys Leu Ser Asp Arg
245 250 255

Leu Gln Tyr Ser Arg Ile Val Gly Gly Trp Asp Leu Leu Pro Arg Ala
260 265 270

Leu Leu Ser Ser Leu Ser Gly Leu Val Leu Leu Asn Ala Pro Val Val
275 280 285

Ala Met Thr Gln Gly Pro His Asp Val His Val Gln Ile Glu Thr Ser
290 295 300

Pro Pro Ala Arg Asn Leu Lys Val Leu Lys Ala Asp Val Val Leu Leu
305 310 315 320

Thr Ala Ser Gly Pro Ala Val Lys Arg Ile Thr Phe Ser Pro Pro Leu
325 330 335

Pro Arg His Met Gln Glu Ala Leu Arg Arg Leu His Tyr Val Pro Ala
340 345 350

Thr Lys Val Phe Leu Ser Phe Arg Arg Pro Phe Trp Arg Glu Glu His
355 360 365

Ile Glu Gly Gly His Ser Asn Thr Asp Arg Pro Ser Arg Met Ile Phe
370 375 380

Tyr Pro Pro Pro Arg Glu Gly Ala Leu Leu Leu Ala Ser Tyr Thr Trp

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385                               390                               395                               400

Ser Asp Ala Ala Ala Ala Phe Ala Gly Leu Ser Arg Glu Glu Ala Leu
      405                               410                               415

Arg Leu Ala Leu Asp Asp Val Ala Ala Leu His Gly Pro Val Val Arg
      420                               425                               430

Gln Leu Trp Asp Gly Thr Gly Val Val Lys Arg Trp Ala Glu Asp Gln
      435                               440                               445

His Ser Gln Gly Gly Phe Val Val Gln Pro Pro Ala Leu Trp Gln Thr
      450                               455                               460

Glu Lys Asp Asp Trp Thr Val Pro Tyr Gly Arg Ile Tyr Phe Ala Gly
465                               470                               475                               480

Glu His Thr Ala Tyr Pro His Gly Trp Val Glu Thr Ala Val Lys Ser
      485                               490                               495

Ala Leu Arg Ala Ala Ile Lys Ile Asn Ser Arg Lys Gly Pro Ala Ser
      500                               505                               510

Asp Thr Ala Ser Pro Glu Gly His Ala Ser Asp Met Glu Gly Gln Gly
      515                               520                               525

His Val His Gly Val Ala Ser Ser Pro Ser His Asp Leu Ala Lys Glu
      530                               535                               540

Glu Gly Ser His Pro Pro Val Gln Gly Gln Leu Ser Leu Gln Asn Thr
545                               550                               555                               560

Thr His Thr Arg Thr Ser His
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<210> 47
<211> 462
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<309> 1986-07-21
<313> (1)..(462)

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<400> 47

Met Ala Arg Val Leu Gly Ala Pro Val Ala Leu Gly Leu Trp Ser Leu
1 5 10 15

Cys Trp Ser Leu Ala Ile Ala Thr Pro Leu Pro Pro Thr Ser Ala His
20 25 30

Gly Asn Val Ala Glu Gly Glu Thr Lys Pro Asp Pro Asp Val Thr Glu
35 40 45

Arg Cys Ser Asp Gly Trp Ser Phe Asp Ala Thr Thr Leu Asp Asp Asn
50 55 60

Gly Thr Met Leu Phe Phe Lys Gly Glu Phe Val Trp Lys Ser His Lys
65 70 75 80

Trp Asp Arg Glu Leu Ile Ser Glu Arg Trp Lys Asn Phe Pro Ser Pro
85 90 95

Val Asp Ala Ala Phe Arg Gln Gly His Asn Ser Val Phe Leu Ile Lys
100 105 110

Gly Asp Lys Val Trp Val Tyr Pro Pro Glu Lys Lys Glu Lys Gly Tyr
115 120 125

Pro Lys Leu Leu Gln Asp Glu Phe Pro Gly Ile Pro Ser Pro Leu Asp
130 135 140

Ala Ala Val Glu Cys His Arg Gly Glu Cys Gln Ala Glu Gly Val Leu
145 150 155 160

Phe Phe Gln Gly Asp Arg Glu Trp Phe Trp Asp Leu Ala Thr Gly Thr
165 170 175

Met Lys Glu Arg Ser Trp Pro Ala Val Gly Asn Cys Ser Ser Ala Leu
180 185 190

Arg Trp Leu Gly Arg Tyr Tyr Cys Phe Gln Gly Asn Gln Phe Leu Arg
195 200 205

Phe Asp Pro Val Arg Gly Glu Val Pro Pro Arg Tyr Pro Arg Asp Val
210 215 220

Arg Asp Tyr Phe Met Pro Cys Pro Gly Arg Gly His Gly His Arg Asn
225 230 235 240

Gly Thr Gly His Gly Asn Ser Thr His His Gly Pro Glu Tyr Met Arg
245 250 255

Cys Ser Pro His Leu Val Leu Ser Ala Leu Thr Ser Asp Asn His Gly
260 265 270

Ala Thr Tyr Ala Phe Ser Gly Thr His Tyr Trp Arg Leu Asp Thr Ser
275 280 285

Arg Asp Gly Trp His Ser Trp Pro Ile Ala His Gln Trp Pro Gln Gly
290 295 300

Pro Ser Ala Val Asp Ala Ala Phe Ser Trp Glu Glu Lys Leu Tyr Leu
305 310 315 320

Val Gln Gly Thr Gln Val Tyr Val Phe Leu Thr Lys Gly Gly Tyr Thr
325 330 335

Leu Val Ser Gly Tyr Pro Lys Arg Leu Glu Lys Glu Val Gly Thr Pro
340 345 350

His Gly Ile Ile Leu Asp Ser Val Asp Ala Ala Phe Ile Cys Pro Gly
355 360 365

Ser Ser Arg Leu His Ile Met Ala Gly Arg Arg Leu Trp Trp Leu Asp
370 375 380

Leu Lys Ser Gly Ala Gln Ala Thr Trp Thr Glu Leu Pro Trp Pro His
385 390 395 400

Glu Lys Val Asp Gly Ala Leu Cys Met Glu Lys Ser Leu Gly Pro Asn
405 410 415

Ser Cys Ser Ala Asn Gly Pro Gly Leu Tyr Leu Ile His Gly Pro Asn
420 425 430

Leu Tyr Cys Tyr Ser Asp Val Glu Lys Leu Asn Ala Ala Lys Ala Leu
435 440 445

Pro Gln Pro Gln Asn Val Thr Ser Leu Leu Gly Cys Thr His
 450 455 460

<210> 48
 <211> 369
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 <213> Homo sapiens

<300>
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 <309> 1996-10-02
 <313> (1)..(369)

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Met Asp Pro Arg Lys Val Asn Glu Leu Arg Ala Phe Val Lys Met Cys
 1 5 10 15

Lys Gln Asp Pro Ser Val Leu His Thr Glu Glu Met Arg Phe Leu Arg
 20 25 30

Glu Trp Val Glu Ser Met Gly Gly Lys Val Pro Pro Ala Thr Gln Lys
 35 40 45

Ala Lys Ser Glu Glu Asn Thr Lys Glu Glu Lys Pro Asp Ser Lys Lys
 50 55 60

Val Glu Glu Asp Leu Lys Ala Asp Glu Pro Ser Ser Glu Glu Ser Asp
 65 70 75 80

Leu Glu Ile Asp Lys Glu Gly Val Ile Glu Pro Asp Thr Asp Ala Pro
 85 90 95

Gln Glu Met Gly Asp Glu Asn Ala Glu Ile Thr Glu Glu Met Met Asp
 100 105 110

Gln Ala Asn Asp Lys Lys Val Ala Ala Ile Glu Ala Leu Asn Asp Gly
 115 120 125

Glu Leu Gln Lys Ala Ile Asp Leu Phe Thr Asp Ala Ile Lys Leu Asn
 130 135 140

Pro Arg Leu Ala Ile Leu Tyr Ala Lys Arg Ala Ser Val Phe Val Lys
 145 150 155 160

Leu Gln Lys Pro Asn Ala Ala Ile Arg Asp Cys Asp Arg Ala Ile Glu

165					170					175					
Ile	Asn	Pro	Asp	Ser	Ala	Gln	Pro	Tyr	Lys	Trp	Arg	Gly	Lys	Ala	His
			180					185					190		
Arg	Leu	Leu	Gly	His	Trp	Glu	Glu	Ala	Ala	His	Asp	Leu	Ala	Leu	Ala
			195				200					205			
Cys	Lys	Leu	Asp	Tyr	Asp	Glu	Asp	Ala	Ser	Ala	Met	Leu	Lys	Glu	Val
			210				215				220				
Gln	Pro	Arg	Ala	Gln	Lys	Ile	Ala	Glu	His	Arg	Arg	Lys	Tyr	Glu	Arg
225					230					235					240
Lys	Arg	Glu	Glu	Arg	Glu	Ile	Lys	Glu	Arg	Ile	Glu	Arg	Val	Lys	Lys
				245					250					255	
Ala	Arg	Glu	Glu	His	Glu	Arg	Ala	Gln	Arg	Glu	Glu	Glu	Ala	Arg	Arg
			260					265					270		
Gln	Ser	Gly	Ala	Gln	Tyr	Gly	Ser	Phe	Pro	Gly	Gly	Phe	Pro	Gly	Gly
			275				280					285			
Met	Pro	Gly	Asn	Phe	Pro	Gly	Gly	Met	Pro	Gly	Met	Gly	Gly	Gly	Met
			290				295				300				
Pro	Gly	Met	Ala	Gly	Met	Pro	Gly	Leu	Asn	Glu	Ile	Leu	Ser	Asp	Pro
305					310					315					320
Glu	Val	Leu	Ala	Ala	Met	Gln	Asp	Pro	Glu	Val	Met	Val	Ala	Phe	Gln
				325					330					335	
Asp	Val	Ala	Gln	Asn	Pro	Ala	Asn	Met	Ser	Lys	Tyr	Gln	Ser	Asn	Pro
			340					345					350		
Lys	Val	Met	Asn	Leu	Ile	Ser	Lys	Leu	Ser	Ala	Lys	Phe	Gly	Gly	Gln
			355				360					365			
Ala															

<210> 49

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<400> 49

Met Gln Leu Met His Ala Asn Ala Gln
1 5

<210> 50
<211> 9
<212> PRT
<213> Homo sapiens

<400> 50

Leu Thr Leu Asp Ser Asn Thr Lys Tyr
1 5

<210> 51
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<400> 51

Phe Val Ile Asp Lys Ser Gly Ser Met
1 5

<210> 52
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<212> PRT
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<400> 52

Tyr Leu Leu Asp Ser Asn Ser Trp Ile
1 5

<210> 53
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<212> PRT
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<400> 53

Tyr Glu Leu Asp Lys Ala Phe Ser Asp
1 5

<210> 54
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<212> PRT
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<400> 54

Ile Lys Ser Gln Gln Ser Glu Val Thr
1 5

<210> 55

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<213> Homo sapiens

<400> 55

Val Gln Ile Glu Thr Ser Pro Pro Ala
1 5

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<211> 9

<212> PRT

<213> Homo sapiens

<400> 56

Ile Ile Leu Asp Ser Val Asp Ala Ala
1 5

<210> 57

<211> 9

<212> PRT

<213> Homo sapiens

<400> 57

Ile Glu Pro Asp Thr Asp Ala Pro Gln
1 5

<210> 58

<211> 17

<212> PRT

<213> Homo sapiens

<400> 58

Asn Ile Gln Pro Ile Phe Ala Val Thr Ser Arg Met Val Lys Thr Tyr
1 5 10 15

Glu

<210> 59

<211> 19

<212> PRT
<213> Homo sapiens

<400> 59

Glu Asn Asn Ile Gln Pro Ile Phe Ala Val Thr Ser Arg Met Val Lys
1 5 10 15

Thr Tyr Glu

<210> 60
<211> 17
<212> PRT
<213> Homo sapiens

<400> 60

Asn Lys Val Phe Gly Glu Asp Ser Val Gly Val Ile Phe Lys Asn Gly
1 5 10 15

Asp

<210> 61
<211> 16
<212> PRT
<213> Homo sapiens

<400> 61

Tyr Pro Glu Gln Leu Lys Met Thr Val Val Lys Leu Ile Ser His Arg
1 5 10 15

<210> 62
<211> 16
<212> PRT
<213> Homo sapiens

<400> 62

Lys Asn Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr
1 5 10 15

<210> 63
<211> 16
<212> PRT
<213> Homo sapiens

<400> 63

Asn Gly Gly His Tyr Thr Tyr Ser Glu Asn Arg Val Glu Lys Asp Gly
1 5 10 15

<210> 64
<211> 15
<212> PRT
<213> Homo sapiens

<400> 64

Gly Pro Asn Asn Tyr Tyr Ser Phe Ala Ser Gln Gln Gln Lys Pro
1 5 10 15

<210> 65
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<212> PRT
<213> Homo sapiens

<400> 65

Gly Pro Asn Asn Tyr Tyr Ser Phe Ala Ser Gln Gln Gln Lys Pro Glu
1 5 10 15

<210> 66
<211> 17
<212> PRT
<213> Homo sapiens

<400> 66

Gly Pro Asn Asn Tyr Tyr Ser Phe Ala Ser Gln Gln Gln Lys Pro Glu
1 5 10 15

Asp

<210> 67
<211> 18
<212> PRT
<213> Homo sapiens

<400> 67

Gly Pro Asn Asn Tyr Tyr Ser Phe Ala Ser Gln Gln Gln Lys Pro Glu
1 5 10 15

Asp Thr

<210> 68

<211> 16
<212> PRT
<213> Homo sapiens

<400> 68

Glu	Lys	Leu	Trp	Phe	Val	Pro	Ala	Lys	Val	Glu	Asp	Ser	Gly	His	Tyr
1				5					10					15	

<210> 69
<211> 14
<212> PRT
<213> Homo sapiens

<400> 69

Ser	Leu	Arg	Glu	Leu	His	Leu	Asp	His	Asn	Gln	Ile	Ser	Arg
1				5					10				

<210> 70
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<212> PRT
<213> Homo sapiens

<400> 70

Leu	Arg	Glu	Leu	His	Leu	Asp	His	Asn	Gln	Ile	Ser	Arg
1				5					10			

<210> 71
<211> 15
<212> PRT
<213> Homo sapiens

<400> 71

Glu	Thr	Met	Lys	Met	Arg	Tyr	Glu	His	Ile	Asp	His	Thr	Phe	Glu
1				5					10				15	

<210> 72
<211> 17
<212> PRT
<213> Homo sapiens

<400> 72

Glu	Thr	Met	Lys	Met	Arg	Tyr	Glu	His	Ile	Asp	His	Thr	Phe	Glu	Ile
1				5					10				15		

Gln

<210> 73
<211> 16
<212> PRT
<213> Homo sapiens

<400> 73

His	Met	Phe	Leu	Gln	Asp	Glu	Ile	Ile	Asp	Lys	Ser	Tyr	Thr	Pro	Ser
1			5						10					15	

<210> 74
<211> 17
<212> PRT
<213> Homo sapiens

<400> 74

Val	Asp	Arg	Tyr	Ile	Ser	Lys	Tyr	Glu	Leu	Asp	Lys	Ala	Phe	Ser	Asp
1			5						10					15	

Arg

<210> 75
<211> 14
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<213> Homo sapiens

<400> 75

Arg	Tyr	Ile	Ser	Lys	Tyr	Glu	Leu	Asp	Lys	Ala	Phe	Ser	Asp
1			5						10				

<210> 76
<211> 19
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<213> Homo sapiens

<400> 76

Leu	Pro	Val	Gly	Arg	Thr	Val	Met	Val	Asn	Ile	Glu	Asn	Pro	Glu	Gly
1			5						10					15	

Ile Pro Val

<210> 77
<211> 20
<212> PRT
<213> Homo sapiens

<400> 77

Leu Pro Val Gly Arg Thr Val Met Val Asn Ile Glu Asn Pro Glu Gly
1 5 10 15

Ile Pro Val Lys
20

<210> 78
<211> 18
<212> PRT
<213> Homo sapiens

<400> 78

Gly Thr Pro His Gly Ile Ile Leu Asp Ser Val Asp Ala Ala Phe Ile
1 5 10 15

Cys Pro

<210> 79
<211> 18
<212> PRT
<213> Homo sapiens

<400> 79

Met Ala Thr Pro Leu Leu Met Gln Ala Leu Pro Met Gly Ala Leu Pro
1 5 10 15

Gln Gly

<210> 80
<211> 17
<212> PRT
<213> Homo sapiens

<400> 80

Met Ala Thr Pro Leu Leu Met Gln Ala Leu Pro Met Gly Ala Leu Pro
1 5 10 15

Gln

<210> 81

<211> 16
<212> PRT
<213> Homo sapiens

<400> 81

Met	Ala	Thr	Pro	Leu	Leu	Met	Gln	Ala	Leu	Pro	Met	Gly	Ala	Leu	Pro
1				5					10					15	

<210> 82
<211> 21
<212> PRT
<213> Homo sapiens

<400> 82

Met	Ala	Thr	Pro	Leu	Leu	Met	Gln	Ala	Leu	Pro	Met	Gly	Ala	Leu	Pro
1				5					10					15	

Gln Gly Pro Met Gln
20

<210> 83
<211> 15
<212> PRT
<213> Homo sapiens

<400> 83

Ala	Thr	Pro	Leu	Leu	Met	Gln	Ala	Leu	Pro	Met	Gly	Ala	Leu	Pro
1				5					10					15

<210> 84
<211> 23
<212> PRT
<213> Homo sapiens

<400> 84

His	Pro	Pro	Val	Gln	Trp	Ala	Phe	Gln	Glu	Thr	Ser	Val	Glu	Ser	Ala
1				5					10					15	

Val Asp Thr Pro Phe Pro Ala
20

<210> 85
<211> 24
<212> PRT
<213> Homo sapiens

<400> 85

His Pro Pro Val Gln Trp Ala Phe Gln Glu Thr Ser Val Glu Ser Ala
1 5 10 15

Val Asp Thr Pro Phe Pro Ala Gly
20

<210> 86
<211> 22
<212> PRT
<213> Homo sapiens

<400> 86

His Pro Pro Val Gln Trp Ala Phe Gln Glu Thr Ser Val Glu Ser Ala
1 5 10 15

Val Asp Thr Pro Phe Pro
20

<210> 87
<211> 17
<212> PRT
<213> Homo sapiens

<400> 87

Lys Ile Tyr Leu Tyr Thr Leu Asn Asp Asn Ala Arg Ser Ser Pro Val
1 5 10 15

Val

<210> 88
<211> 16
<212> PRT
<213> Homo sapiens

<400> 88

Lys Ile Tyr Leu Tyr Thr Leu Asn Asp Asn Ala Arg Ser Ser Pro Val
1 5 10 15

<210> 89
<211> 17
<212> PRT
<213> Homo sapiens

<400> 89

Ile Tyr Leu Tyr Thr Leu Asn Asp Asn Ala Arg Ser Ser Pro Val Val
1 5 10 15

Ile

<210> 90
<211> 15
<212> PRT
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<400> 90

Ile Tyr Leu Tyr Thr Leu Asn Asp Asn Ala Arg Ser Ser Pro Val
1 5 10 15

<210> 91
<211> 16
<212> PRT
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<400> 91

Ile Tyr Leu Tyr Thr Leu Asn Asp Asn Ala Arg Ser Ser Pro Val Val
1 5 10 15

<210> 92
<211> 15
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<400> 92

Tyr Asn Ser Tyr Ser Val Ser Asn Ser Glu Lys Asp Ile Met Ala
1 5 10 15

<210> 93
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<400> 93

Ala Gly Ser Leu Thr Leu Ser Lys Thr Glu Leu Gly Lys Lys Ala
1 5 10 15

<210> 94
<211> 16
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<400> 94

Ala Gly Ser Leu Thr Leu Ser Lys Thr Glu Leu Gly Lys Lys Ala Asp
1 5 10 15

<210> 95

<211> 14

<212> PRT

<213> Homo sapiens

<400> 95

Val Pro Lys Asp Tyr Thr Gly Glu Asp Val Thr Pro Gln Asn
1 5 10

<210> 96

<211> 13

<212> PRT

<213> Homo sapiens

<400> 96

Asp Ser Lys Phe His Gln Ala Ile Asn Asp Ala His Gln
1 5 10

<210> 97

<211> 15

<212> PRT

<213> Homo sapiens

<400> 97

Met Pro Leu Glu Phe Lys Thr Leu Asn Val Leu His Asn Arg Gly
1 5 10 15

<210> 98

<211> 15

<212> PRT

<213> Homo sapiens

<400> 98

Ala Thr Arg Ser Ile Gln Val Asp Gly Lys Thr Ile Lys Ala Gln
1 5 10 15

<210> 99

<211> 18

<212> PRT

<213> Homo sapiens

<400> 99

Ala Thr Arg Ser Ile Gln Val Asp Gly Lys Thr Ile Lys Ala Gln Ile
1 5 10 15

Trp Asp

<210> 100
<211> 14
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<400> 100

Thr Arg Ser Ile Gln Val Asp Gly Lys Thr Ile Lys Ala Gln
1 5 10

<210> 101
<211> 17
<212> PRT
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<400> 101

Ala Thr Arg Ser Ile Gln Val Asp Gly Lys Thr Ile Lys Ala Gln Ile
1 5 10 15

Trp

<210> 102
<211> 13
<212> PRT
<213> Homo sapiens

<400> 102

Arg Ser Ile Gln Val Asp Gly Lys Thr Ile Lys Ala Gln
1 5 10

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<212> PRT
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<400> 103

Ile Gln Pro Ile Phe Ala Val Thr Ser
1 5

<210> 104

<211> 9
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Val Phe Gly Glu Asp Ser Val Gly Val
1 5

<210> 105
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<400> 105

Leu Lys Met Thr Val Val Lys Leu Ile
1 5

<210> 106
<211> 9
<212> PRT
<213> Homo sapiens

<400> 106

Tyr Leu Gln Met Asn Ser Leu Arg Ala
1 5

<210> 107
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<400> 107

Tyr Thr Tyr Ser Glu Asn Arg Val Glu
1 5

<210> 108
<211> 9
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Tyr Tyr Ser Phe Ala Ser Gln Gln Gln
1 5

<210> 109
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Phe Val Pro Ala Lys Val Glu Asp Ser
1 5

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<400> 110

Leu His Leu Asp His Asn Gln Ile Ser
1 5

<210> 111

<211> 9

<212> PRT

<213> Homo sapiens

<400> 111

Met Arg Tyr Glu His Ile Asp His Thr
1 5

<210> 112

<211> 9

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<213> Homo sapiens

<400> 112

Phe Leu Gln Asp Glu Ile Ile Asp Lys
1 5

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<212> PRT

<213> Homo sapiens

<400> 113

Met Val Asn Ile Glu Asn Pro Glu Gly
1 5

<210> 114

<211> 9

<212> PRT

<213> Homo sapiens

<400> 114

Leu Met Gln Ala Leu Pro Met Gly Ala
1 5

<210> 115

<211> 9

<212> PRT

<213> Homo sapiens

<400> 115

Trp Ala Phe Gln Glu Thr Ser Val Glu
1 5

<210> 116

<211> 9

<212> PRT

<213> Homo sapiens

<400> 116

Tyr Thr Leu Asn Asp Asn Ala Arg Ser
1 5

<210> 117

<211> 9

<212> PRT

<213> Homo sapiens

<400> 117

Tyr Ser Val Ser Asn Ser Glu Lys Asp
1 5

<210> 118

<211> 9

<212> PRT

<213> Homo sapiens

<400> 118

Leu Thr Leu Ser Lys Thr Glu Leu Gly
1 5

<210> 119

<211> 9

<212> PRT

<213> Homo sapiens

<400> 119

Tyr Thr Gly Glu Asp Val Thr Pro Gln
1 5

<210> 120
<211> 9
<212> PRT
<213> Homo sapiens

<400> 120

Phe His Gln Ala Ile Asn Asp Ala His
1 5

<210> 121
<211> 9
<212> PRT
<213> Homo sapiens

<400> 121

Phe Lys Thr Leu Asn Val Leu His Asn
1 5

<210> 122
<211> 9
<212> PRT
<213> Homo sapiens

<400> 122

Ile Gln Val Asp Gly Lys Thr Ile Lys
1 5

<210> 123
<211> 769
<212> PRT
<213> Homo sapiens

<300>
<308> Swissprot/P05107
<309> 1987-08-13
<313> (1)..(769)

<400> 123

Met Leu Gly Leu Arg Pro Pro Leu Leu Ala Leu Val Gly Leu Leu Ser
1 5 10 15

Leu Gly Cys Val Leu Ser Gln Glu Cys Thr Lys Phe Lys Val Ser Ser
20 25 30

Cys	Arg	Glu	Cys	Ile	Glu	Ser	Gly	Pro	Gly	Cys	Thr	Trp	Cys	Gln	Lys		
		35					40					45					
Leu	Asn	Phe	Thr	Gly	Pro	Gly	Asp	Pro	Asp	Ser	Ile	Arg	Cys	Asp	Thr		
	50					55					60						
Arg	Pro	Gln	Leu	Leu	Met	Arg	Gly	Cys	Ala	Ala	Asp	Asp	Ile	Met	Asp		
65					70					75					80		
Pro	Thr	Ser	Leu	Ala	Glu	Thr	Gln	Glu	Asp	His	Asn	Gly	Gly	Gln	Lys		
				85					90					95			
Gln	Leu	Ser	Pro	Gln	Lys	Val	Thr	Leu	Tyr	Leu	Arg	Pro	Gly	Gln	Ala		
			100					105					110				
Ala	Ala	Phe	Asn	Val	Thr	Phe	Arg	Arg	Ala	Lys	Gly	Tyr	Pro	Ile	Asp		
		115					120					125					
Leu	Tyr	Tyr	Leu	Met	Asp	Leu	Ser	Tyr	Ser	Met	Leu	Asp	Asp	Leu	Arg		
	130					135					140						
Asn	Val	Lys	Lys	Leu	Gly	Gly	Asp	Leu	Leu	Arg	Ala	Leu	Asn	Glu	Ile		
145					150					155					160		
Thr	Glu	Ser	Gly	Arg	Ile	Gly	Phe	Gly	Ser	Phe	Val	Asp	Lys	Thr	Val		
				165					170					175			
Leu	Pro	Phe	Val	Asn	Thr	His	Pro	Asp	Lys	Leu	Arg	Asn	Pro	Cys	Pro		
			180					185					190				
Asn	Lys	Glu	Lys	Glu	Cys	Gln	Pro	Pro	Phe	Ala	Phe	Arg	His	Val	Leu		
		195					200					205					
Lys	Leu	Thr	Asn	Asn	Ser	Asn	Gln	Phe	Gln	Thr	Glu	Val	Gly	Lys	Gln		
	210					215					220						
Leu	Ile	Ser	Gly	Asn	Leu	Asp	Ala	Pro	Glu	Gly	Gly	Leu	Asp	Ala	Met		
225					230					235					240		
Met	Gln	Val	Ala	Ala	Cys	Pro	Glu	Glu	Ile	Gly	Trp	Arg	Asn	Val	Thr		
				245					250					255			
Arg	Leu	Leu	Val	Phe	Ala	Thr	Asp	Asp	Gly	Phe	His	Phe	Ala	Gly	Asp		

260	265	270
Gly Lys Leu Gly Ala Ile Leu Thr Pro Asn Asp Gly Arg Cys His Leu		
275	280	285
Glu Asp Asn Leu Tyr Lys Arg Ser Asn Glu Phe Asp Tyr Pro Ser Val		
290	295	300
Gly Gln Leu Ala His Lys Leu Ala Glu Asn Asn Ile Gln Pro Ile Phe		
305	310	315
Ala Val Thr Ser Arg Met Val Lys Thr Tyr Glu Lys Leu Thr Glu Ile		
	325	330
Ile Pro Lys Ser Ala Val Gly Glu Leu Ser Glu Asp Ser Ser Asn Val		
	340	345
Val His Leu Ile Lys Asn Ala Tyr Asn Lys Leu Ser Ser Arg Val Phe		
	355	360
Leu Asp His Asn Ala Leu Pro Asp Thr Leu Lys Val Thr Tyr Asp Ser		
	370	375
Phe Cys Ser Asn Gly Val Thr His Arg Asn Gln Pro Arg Gly Asp Cys		
385	390	395
Asp Gly Val Gln Ile Asn Val Pro Ile Thr Phe Gln Val Lys Val Thr		
	405	410
Ala Thr Glu Cys Ile Gln Glu Gln Ser Phe Val Ile Arg Ala Leu Gly		
	420	425
Phe Thr Asp Ile Val Thr Val Gln Val Leu Pro Gln Cys Glu Cys Arg		
	435	440
Cys Arg Asp Gln Ser Arg Asp Arg Ser Leu Cys His Gly Lys Gly Phe		
	450	455
Leu Glu Cys Gly Ile Cys Arg Cys Asp Thr Gly Tyr Ile Gly Lys Asn		
465	470	475
Cys Glu Cys Gln Thr Gln Gly Arg Ser Ser Gln Glu Leu Glu Gly Ser		
	485	490
		495

Cys Arg Lys Asp Asn Asn Ser Ile Ile Cys Ser Gly Leu Gly Asp Cys
500 505 510

Val Cys Gly Gln Cys Leu Cys His Thr Ser Asp Val Pro Gly Lys Leu
515 520 525

Ile Tyr Gly Gln Tyr Cys Glu Cys Asp Thr Ile Asn Cys Glu Arg Tyr
530 535 540

Asn Gly Gln Val Cys Gly Gly Pro Gly Arg Gly Leu Cys Phe Cys Gly
545 550 555 560

Lys Cys Arg Cys His Pro Gly Phe Glu Gly Ser Ala Cys Gln Cys Glu
565 570 575

Arg Thr Thr Glu Gly Cys Leu Asn Pro Arg Arg Val Glu Cys Ser Gly
580 585 590

Arg Gly Arg Cys Arg Cys Asn Val Cys Glu Cys His Ser Gly Tyr Gln
595 600 605

Leu Pro Leu Cys Gln Glu Cys Pro Gly Cys Pro Ser Pro Cys Gly Lys
610 615 620

Tyr Ile Ser Cys Ala Glu Cys Leu Lys Phe Glu Lys Gly Pro Phe Gly
625 630 635 640

Lys Asn Cys Ser Ala Ala Cys Pro Gly Leu Gln Leu Ser Asn Asn Pro
645 650 655

Val Lys Gly Arg Thr Cys Lys Glu Arg Asp Ser Glu Gly Cys Trp Val
660 665 670

Ala Tyr Thr Leu Glu Gln Gln Asp Gly Met Asp Arg Tyr Leu Ile Tyr
675 680 685

Val Asp Glu Ser Arg Glu Cys Val Ala Gly Pro Asn Ile Ala Ala Ile
690 695 700

Val Gly Gly Thr Val Ala Gly Ile Val Leu Ile Gly Ile Leu Leu Leu
705 710 715 720

Val Ile Trp Lys Ala Leu Ile His Leu Ser Asp Leu Arg Glu Tyr Arg
725 730 735

Arg Phe Glu Lys Glu Lys Leu Lys Ser Gln Trp Asn Asn Asp Asn Pro
740 745 750

Leu Phe Lys Ser Ala Thr Thr Thr Val Met Asn Pro Lys Phe Ala Glu
755 760 765

Ser

<210> 124
<211> 1070
<212> PRT
<213> Homo sapiens

<300>
<308> Swissprot/P42338
<309> 1995-11-01
<313> (1)..(1070)

<400> 124

Met Cys Phe Ser Phe Ile Met Pro Pro Ala Met Ala Asp Ile Leu Asp
1 5 10 15

Ile Trp Ala Val Asp Ser Gln Ile Ala Ser Asp Gly Ser Ile Pro Val
20 25 30

Asp Phe Leu Leu Pro Thr Gly Ile Tyr Ile Gln Leu Glu Val Pro Arg
35 40 45

Glu Ala Thr Ile Ser Tyr Ile Lys Gln Met Leu Trp Lys Gln Val His
50 55 60

Asn Tyr Pro Met Phe Asn Leu Leu Met Asp Ile Asp Ser Tyr Met Phe
65 70 75 80

Ala Cys Val Asn Gln Thr Ala Val Tyr Glu Glu Leu Glu Asp Glu Thr
85 90 95

Arg Arg Leu Cys Asp Val Arg Pro Phe Leu Pro Val Leu Lys Leu Val
100 105 110

Thr Arg Ser Cys Asp Pro Gly Glu Lys Leu Asp Ser Lys Ile Gly Val
115 120 125

Leu Ile Gly Lys Gly Leu His Glu Phe Asp Ser Leu Lys Asp Pro Glu
130 135 140

Val Asn Glu Phe Arg Arg Lys Met Arg Lys Phe Ser Glu Glu Lys Ile
145 150 155 160

Leu Ser Leu Val Gly Leu Ser Trp Met Asp Trp Leu Lys Gln Thr Tyr
165 170 175

Pro Pro Glu His Glu Pro Ser Ile Pro Glu Asn Leu Glu Asp Lys Leu
180 185 190

Tyr Gly Gly Lys Leu Ile Val Ala Val His Phe Glu Asn Cys Gln Asp
195 200 205

Val Phe Ser Phe Gln Val Ser Pro Asn Met Asn Pro Ile Lys Val Asn
210 215 220

Glu Leu Ala Ile Gln Lys Arg Leu Thr Ile His Gly Lys Glu Asp Glu
225 230 235 240

Val Ser Pro Tyr Asp Tyr Val Leu Gln Val Ser Gly Arg Val Glu Tyr
245 250 255

Val Phe Gly Asp His Pro Leu Ile Gln Phe Gln Tyr Ile Arg Asn Cys
260 265 270

Val Met Asn Arg Ala Leu Pro His Phe Ile Leu Val Glu Cys Cys Lys
275 280 285

Ile Lys Lys Met Tyr Glu Gln Glu Met Ile Ala Ile Glu Ala Ala Ile
290 295 300

Asn Arg Asn Ser Ser Asn Leu Pro Leu Pro Leu Pro Pro Lys Lys Thr
305 310 315 320

Arg Ile Ile Ser His Val Trp Glu Asn Asn Asn Pro Phe Gln Ile Val
325 330 335

Leu Val Lys Gly Asn Lys Leu Asn Thr Glu Glu Thr Val Lys Val His

340	345	350
Val Arg Ala Gly Leu Phe His Gly Thr Glu Leu Leu Cys Lys Thr Ile 355 360 365		
Val Ser Ser Glu Val Ser Gly Lys Asn Asp His Ile Trp Asn Glu Pro 370 375 380		
Leu Glu Phe Asp Ile Asn Ile Cys Asp Leu Pro Arg Met Ala Arg Leu 385 390 395 400		
Cys Phe Ala Val Tyr Ala Val Leu Asp Lys Val Lys Thr Lys Lys Ser 405 410 415		
Thr Lys Thr Ile Asn Pro Ser Lys Tyr Gln Thr Ile Arg Lys Ala Gly 420 425 430		
Lys Val His Tyr Pro Val Ala Trp Val Asn Thr Met Val Phe Asp Phe 435 440 445		
Lys Gly Gln Leu Arg Thr Gly Asp Ile Ile Leu His Ser Trp Ser Ser 450 455 460		
Phe Pro Asp Glu Leu Glu Glu Met Leu Asn Pro Met Gly Thr Val Gln 465 470 475 480		
Thr Asn Pro Tyr Thr Glu Asn Ala Thr Ala Leu His Val Lys Phe Pro 485 490 495		
Glu Asn Lys Lys Gln Pro Tyr Tyr Tyr Pro Pro Phe Asp Lys Ile Ile 500 505 510		
Glu Lys Ala Ala Glu Ile Ala Ser Ser Asp Ser Ala Asn Val Ser Ser 515 520 525		
Arg Gly Gly Lys Lys Phe Leu Pro Val Leu Lys Glu Ile Leu Asp Arg 530 535 540		
Asp Pro Leu Ser Gln Leu Cys Glu Asn Glu Met Asp Leu Ile Trp Thr 545 550 555 560		
Leu Arg Gln Asp Cys Arg Glu Ile Phe Pro Gln Ser Leu Pro Lys Leu 565 570 575		

Leu Leu Ser Ile Lys Trp Asn Lys Leu Glu Asp Val Ala Gln Leu Gln
580 585 590

Ala Leu Leu Gln Ile Trp Pro Lys Leu Pro Pro Arg Glu Ala Leu Glu
595 600 605

Leu Leu Asp Phe Asn Tyr Pro Asp Gln Tyr Val Arg Glu Tyr Ala Val
610 615 620

Gly Cys Leu Arg Gln Met Ser Asp Glu Glu Leu Ser Gln Tyr Leu Leu
625 630 635 640

Gln Leu Val Gln Val Leu Lys Tyr Glu Pro Phe Leu Asp Cys Ala Leu
645 650 655

Ser Arg Phe Leu Leu Glu Arg Ala Leu Gly Asn Arg Arg Ile Gly Gln
660 665 670

Phe Leu Phe Trp His Leu Arg Ser Glu Val His Ile Pro Ala Val Ser
675 680 685

Val Gln Phe Gly Val Ile Leu Glu Ala Tyr Cys Arg Gly Ser Val Gly
690 695 700

His Met Lys Val Leu Ser Lys Gln Val Glu Ala Leu Asn Lys Leu Lys
705 710 715 720

Thr Leu Asn Ser Leu Ile Lys Leu Asn Ala Val Lys Leu Asn Arg Ala
725 730 735

Lys Gly Lys Glu Ala Met His Thr Cys Leu Lys Gln Ser Ala Tyr Arg
740 745 750

Glu Ala Leu Ser Asp Leu Gln Ser Pro Leu Asn Pro Cys Val Ile Leu
755 760 765

Ser Glu Leu Tyr Val Glu Lys Cys Lys Tyr Met Asp Ser Lys Met Lys
770 775 780

Pro Leu Trp Leu Val Tyr Asn Asn Lys Val Phe Gly Glu Asp Ser Val
785 790 795 800

Gly Val Ile Phe Lys Asn Gly Asp Asp Leu Arg Gln Asp Met Leu Thr
805 810 815

Leu Gln Met Leu Arg Leu Met Asp Leu Leu Trp Lys Glu Ala Gly Leu
820 825 830

Asp Leu Arg Met Leu Pro Tyr Gly Cys Leu Ala Thr Gly Asp Arg Ser
835 840 845

Gly Leu Ile Glu Val Val Ser Thr Ser Glu Thr Ile Ala Asp Ile Gln
850 855 860

Leu Asn Ser Ser Asn Val Ala Ala Ala Ala Ala Phe Asn Lys Asp Ala
865 870 875 880

Leu Leu Asn Trp Leu Lys Glu Tyr Asn Ser Gly Asp Asp Leu Asp Arg
885 890 895

Ala Ile Glu Glu Phe Thr Leu Ser Cys Ala Gly Tyr Cys Val Ala Ser
900 905 910

Tyr Val Leu Gly Ile Gly Asp Arg His Ser Asp Asn Ile Met Val Lys
915 920 925

Lys Thr Gly Gln Leu Phe His Ile Asp Phe Gly His Ile Leu Gly Asn
930 935 940

Phe Lys Ser Lys Phe Gly Ile Lys Arg Glu Arg Val Pro Phe Ile Leu
945 950 955 960

Thr Tyr Asp Phe Ile His Val Ile Gln Gln Gly Lys Thr Gly Asn Thr
965 970 975

Glu Lys Phe Gly Arg Phe Arg Gln Cys Cys Glu Asp Ala Tyr Leu Ile
980 985 990

Leu Arg Arg His Gly Asn Leu Phe Ile Thr Leu Phe Ala Leu Met Leu
995 1000 1005

Thr Ala Gly Leu Pro Glu Leu Thr Ser Val Lys Asp Ile Gln Tyr
1010 1015 1020

Leu Lys Asp Ser Leu Ala Leu Gly Lys Ser Glu Glu Glu Ala Leu
 1025 1030 1035

Lys Gln Phe Lys Gln Lys Phe Asp Glu Ala Leu Arg Glu Ser Trp
 1040 1045 1050

Thr Thr Lys Val Asn Trp Met Ala His Thr Val Arg Lys Asp Tyr
 1055 1060 1065

Arg Ser
 1070

<210> 125
 <211> 431
 <212> PRT
 <213> Homo sapiens

<300>
 <308> Swissprot/P00749
 <309> 1986-07-21
 <313> (1)..(431)

<400> 125

Met Arg Ala Leu Leu Ala Arg Leu Leu Leu Cys Val Leu Val Val Ser
 1 5 10 15

Asp Ser Lys Gly Ser Asn Glu Leu His Gln Val Pro Ser Asn Cys Asp
 20 25 30

Cys Leu Asn Gly Gly Thr Cys Val Ser Asn Lys Tyr Phe Ser Asn Ile
 35 40 45

His Trp Cys Asn Cys Pro Lys Lys Phe Gly Gly Gln His Cys Glu Ile
 50 55 60

Asp Lys Ser Lys Thr Cys Tyr Glu Gly Asn Gly His Phe Tyr Arg Gly
 65 70 75 80

Lys Ala Ser Thr Asp Thr Met Gly Arg Pro Cys Leu Pro Trp Asn Ser
 85 90 95

Ala Thr Val Leu Gln Gln Thr Tyr His Ala His Arg Ser Asp Ala Leu
 100 105 110

Gln Leu Gly Leu Gly Lys His Asn Tyr Cys Arg Asn Pro Asp Asn Arg

115							120						125			
Arg	Arg	Pro	Trp	Cys	Tyr	Val	Gln	Val	Gly	Leu	Lys	Pro	Leu	Val	Gln	
130						135					140					
Glu	Cys	Met	Val	His	Asp	Cys	Ala	Asp	Gly	Lys	Lys	Pro	Ser	Ser	Pro	
145					150					155					160	
Pro	Glu	Glu	Leu	Lys	Phe	Gln	Cys	Gly	Gln	Lys	Thr	Leu	Arg	Pro	Arg	
				165					170					175		
Phe	Lys	Ile	Ile	Gly	Gly	Glu	Phe	Thr	Thr	Ile	Glu	Asn	Gln	Pro	Trp	
			180					185					190			
Phe	Ala	Ala	Ile	Tyr	Arg	Arg	His	Arg	Gly	Gly	Ser	Val	Thr	Tyr	Val	
	195						200					205				
Cys	Gly	Gly	Ser	Leu	Met	Ser	Pro	Cys	Trp	Val	Ile	Ser	Ala	Thr	His	
210						215					220					
Cys	Phe	Ile	Asp	Tyr	Pro	Lys	Lys	Glu	Asp	Tyr	Ile	Val	Tyr	Leu	Gly	
225					230					235					240	
Arg	Ser	Arg	Leu	Asn	Ser	Asn	Thr	Gln	Gly	Glu	Met	Lys	Phe	Glu	Val	
				245					250					255		
Glu	Asn	Leu	Ile	Leu	His	Lys	Asp	Tyr	Ser	Ala	Asp	Thr	Leu	Ala	His	
		260						265					270			
His	Asn	Asp	Ile	Ala	Leu	Leu	Lys	Ile	Arg	Ser	Lys	Glu	Gly	Arg	Cys	
		275					280					285				
Ala	Gln	Pro	Ser	Arg	Thr	Ile	Gln	Thr	Ile	Cys	Leu	Pro	Ser	Met	Tyr	
	290					295					300					
Asn	Asp	Pro	Gln	Phe	Gly	Thr	Ser	Cys	Glu	Ile	Thr	Gly	Phe	Gly	Lys	
305					310					315					320	
Glu	Asn	Ser	Thr	Asp	Tyr	Leu	Tyr	Pro	Glu	Gln	Leu	Lys	Met	Thr	Val	
				325					330					335		
Val	Lys	Leu	Ile	Ser	His	Arg	Glu	Cys	Gln	Gln	Pro	His	Tyr	Tyr	Gly	
		340					345						350			

Ser Glu Val Thr Thr Lys Met Leu Cys Ala Ala Asp Pro Gln Trp Lys
355 360 365

Thr Asp Ser Cys Gln Gly Asp Ser Gly Gly Pro Leu Val Cys Ser Leu
370 375 380

Gln Gly Arg Met Thr Leu Thr Gly Ile Val Ser Trp Gly Arg Gly Cys
385 390 395 400

Ala Leu Lys Asp Lys Pro Gly Val Tyr Thr Arg Val Ser His Phe Leu
405 410 415

Pro Trp Ile Arg Ser His Thr Lys Glu Glu Asn Gly Leu Ala Leu
420 425 430

<210> 126
<211> 117
<212> PRT
<213> Homo sapiens

<300>
<308> Swissprot/P01764
<309> 1986-07-21
<313> (1)..(117)

<400> 126

Met Glu Phe Gly Leu Ser Trp Leu Phe Leu Val Ala Ile Leu Lys Gly
1 5 10 15

Val Gln Cys Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln
20 25 30

Pro Gly Gly Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe
35 40 45

Ser Ser Tyr Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu
50 55 60

Glu Trp Val Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Gly
65 70 75 80

Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn
85 90 95

Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val
100 105 110

Tyr Tyr Cys Ala Lys
115

<210> 127
<211> 189
<212> PRT
<213> Homo sapiens

<300>
<308> Swissprot/Q99497
<309> 1997-05-01
<313> (1)..(189)

<400> 127

Met Ala Ser Lys Arg Ala Leu Val Ile Leu Ala Lys Gly Ala Glu Glu
1 5 10 15

Met Glu Thr Val Ile Pro Val Asp Val Met Arg Arg Ala Gly Ile Lys
20 25 30

Val Thr Val Ala Gly Leu Ala Gly Lys Asp Pro Val Gln Cys Ser Arg
35 40 45

Asp Val Val Ile Cys Pro Asp Ala Ser Leu Glu Asp Ala Lys Lys Glu
50 55 60

Gly Pro Tyr Asp Val Val Val Leu Pro Gly Gly Asn Leu Gly Ala Gln
65 70 75 80

Asn Leu Ser Glu Ser Ala Ala Val Lys Glu Ile Leu Lys Glu Gln Glu
85 90 95

Asn Arg Lys Gly Leu Ile Ala Ala Ile Cys Ala Gly Pro Thr Ala Leu
100 105 110

Leu Ala His Glu Ile Gly Phe Gly Ser Lys Val Thr Thr His Pro Leu
115 120 125

Ala Lys Asp Lys Met Met Asn Gly Gly His Tyr Thr Tyr Ser Glu Asn
130 135 140

Arg Val Glu Lys Asp Gly Leu Ile Leu Thr Ser Arg Gly Pro Gly Thr
145 150 155 160

Ser Phe Glu Phe Ala Leu Ala Ile Val Glu Ala Leu Asn Gly Lys Glu
165 170 175

Val Ala Ala Gln Val Lys Ala Pro Leu Val Leu Lys Asp
180 185

<210> 128
<211> 257
<212> PRT
<213> Homo sapiens

<300>
<308> Swissprot/P48556
<309> 1996-02-01
<313> (1)..(257)

<400> 128

Met Tyr Glu Gln Leu Lys Gly Glu Trp Asn Arg Lys Ser Pro Asn Leu
1 5 10 15

Ser Lys Cys Gly Glu Glu Leu Gly Arg Leu Lys Leu Val Leu Leu Glu
20 25 30

Leu Asn Phe Leu Pro Thr Thr Gly Thr Lys Leu Thr Lys Gln Gln Leu
35 40 45

Ile Leu Ala Arg Asp Ile Leu Glu Ile Gly Ala Gln Trp Ser Ile Leu
50 55 60

Arg Lys Asp Ile Pro Ser Phe Glu Arg Tyr Met Ala Gln Leu Lys Cys
65 70 75 80

Tyr Tyr Phe Asp Tyr Lys Glu Gln Leu Pro Glu Ser Ala Tyr Met His
85 90 95

Gln Leu Leu Gly Leu Asn Leu Leu Phe Leu Leu Ser Gln Asn Arg Val
100 105 110

Ala Glu Phe His Thr Glu Leu Glu Arg Leu Pro Ala Lys Asp Ile Gln
115 120 125

Thr Asn Val Tyr Ile Lys His Pro Val Ser Leu Glu Gln Tyr Leu Met

130	135	140
Glu Gly Ser Tyr Asn Lys Val Phe Leu Ala Lys Gly Asn Ile Pro Ala		
145	150	155 160
Glu Ser Tyr Thr Phe Phe Ile Asp Ile Leu Leu Asp Thr Ile Arg Asp		
	165	170 175
Glu Ile Ala Gly Cys Ile Glu Lys Ala Tyr Glu Lys Ile Leu Phe Thr		
	180	185 190
Glu Ala Thr Arg Ile Leu Phe Phe Asn Thr Pro Lys Lys Met Thr Asp		
	195	200 205
Tyr Ala Lys Lys Arg Gly Trp Val Leu Gly Pro Asn Asn Tyr Tyr Ser		
	210	215 220
Phe Ala Ser Gln Gln Gln Lys Pro Glu Asp Thr Thr Ile Pro Ser Thr		
225	230	235 240
Glu Leu Ala Lys Gln Val Ile Glu Tyr Ala Arg Gln Leu Glu Met Ile		
	245	250 255

Val

<210> 129
 <211> 569
 <212> PRT
 <213> Homo sapiens

<300>
 <308> Swissprot/P14778
 <309> 1990-04-01
 <313> (1)..(569)

<400> 129

Met Lys Val Leu Leu Arg Leu Ile Cys Phe Ile Ala Leu Leu Ile Ser
1 5 10 15
Ser Leu Glu Ala Asp Lys Cys Lys Glu Arg Glu Glu Lys Ile Ile Leu
20 25 30
Val Ser Ser Ala Asn Glu Ile Asp Val Arg Pro Cys Pro Leu Asn Pro
35 40 45

Asn Glu His Lys Gly Thr Ile Thr Trp Tyr Lys Asp Asp Ser Lys Thr
50 55 60

Pro Val Ser Thr Glu Gln Ala Ser Arg Ile His Gln His Lys Glu Lys
65 70 75 80

Leu Trp Phe Val Pro Ala Lys Val Glu Asp Ser Gly His Tyr Tyr Cys
85 90 95

Val Val Arg Asn Ser Ser Tyr Cys Leu Arg Ile Lys Ile Ser Ala Lys
100 105 110

Phe Val Glu Asn Glu Pro Asn Leu Cys Tyr Asn Ala Gln Ala Ile Phe
115 120 125

Lys Gln Lys Leu Pro Val Ala Gly Asp Gly Gly Leu Val Cys Pro Tyr
130 135 140

Met Glu Phe Phe Lys Asn Glu Asn Asn Glu Leu Pro Lys Leu Gln Trp
145 150 155 160

Tyr Lys Asp Cys Lys Pro Leu Leu Leu Asp Asn Ile His Phe Ser Gly
165 170 175

Val Lys Asp Arg Leu Ile Val Met Asn Val Ala Glu Lys His Arg Gly
180 185 190

Asn Tyr Thr Cys His Ala Ser Tyr Thr Tyr Leu Gly Lys Gln Tyr Pro
195 200 205

Ile Thr Arg Val Ile Glu Phe Ile Thr Leu Glu Glu Asn Lys Pro Thr
210 215 220

Arg Pro Val Ile Val Ser Pro Ala Asn Glu Thr Met Glu Val Asp Leu
225 230 235 240

Gly Ser Gln Ile Gln Leu Ile Cys Asn Val Thr Gly Gln Leu Ser Asp
245 250 255

Ile Ala Tyr Trp Lys Trp Asn Gly Ser Val Ile Asp Glu Asp Asp Pro
260 265 270

Val Leu Gly Glu Asp Tyr Tyr Ser Val Glu Asn Pro Ala Asn Lys Arg
275 280 285

Arg Ser Thr Leu Ile Thr Val Leu Asn Ile Ser Glu Ile Glu Ser Arg
290 295 300

Phe Tyr Lys His Pro Phe Thr Cys Phe Ala Lys Asn Thr His Gly Ile
305 310 315 320

Asp Ala Ala Tyr Ile Gln Leu Ile Tyr Pro Val Thr Asn Phe Gln Lys
325 330 335

His Met Ile Gly Ile Cys Val Thr Leu Thr Val Ile Ile Val Cys Ser
340 345 350

Val Phe Ile Tyr Lys Ile Phe Lys Ile Asp Ile Val Leu Trp Tyr Arg
355 360 365

Asp Ser Cys Tyr Asp Phe Leu Pro Ile Lys Ala Ser Asp Gly Lys Thr
370 375 380

Tyr Asp Ala Tyr Ile Leu Tyr Pro Lys Thr Val Gly Glu Gly Ser Thr
385 390 395 400

Ser Asp Cys Asp Ile Phe Val Phe Lys Val Leu Pro Glu Val Leu Glu
405 410 415

Lys Gln Cys Gly Tyr Lys Leu Phe Ile Tyr Gly Arg Asp Asp Tyr Val
420 425 430

Gly Glu Asp Ile Val Glu Val Ile Asn Glu Asn Val Lys Lys Ser Arg
435 440 445

Arg Leu Ile Ile Ile Leu Val Arg Glu Thr Ser Gly Phe Ser Trp Leu
450 455 460

Gly Gly Ser Ser Glu Glu Gln Ile Ala Met Tyr Asn Ala Leu Val Gln
465 470 475 480

Asp Gly Ile Lys Val Val Leu Leu Glu Leu Glu Lys Ile Gln Asp Tyr
485 490 495

Glu Lys Met Pro Glu Ser Ile Lys Phe Ile Lys Gln Lys His Gly Ala
500 505 510

Ile Arg Trp Ser Gly Asp Phe Thr Gln Gly Pro Gln Ser Ala Lys Thr
515 520 525

Arg Phe Trp Lys Asn Val Arg Tyr His Met Pro Val Gln Arg Arg Ser
530 535 540

Pro Ser Ser Lys His Gln Leu Leu Ser Pro Ala Thr Lys Glu Lys Leu
545 550 555 560

Gln Arg Glu Ala His Val Pro Leu Gly
565

<210> 130
<211> 376
<212> PRT
<213> Homo sapiens

<300>
<308> Swissprot/Q06828
<309> 1994-06-01
<313> (1)..(376)

<400> 130

Met Gln Trp Thr Ser Leu Leu Leu Leu Ala Gly Leu Phe Ser Leu Ser
1 5 10 15

Gln Ala Gln Tyr Glu Asp Asp Pro His Trp Trp Phe His Tyr Leu Arg
20 25 30

Ser Gln Gln Ser Thr Tyr Tyr Asp Pro Tyr Asp Pro Tyr Pro Tyr Glu
35 40 45

Thr Tyr Glu Pro Tyr Pro Tyr Gly Val Asp Glu Gly Pro Ala Tyr Thr
50 55 60

Tyr Gly Ser Pro Ser Pro Pro Asp Pro Arg Asp Cys Pro Gln Glu Cys
65 70 75 80

Asp Cys Pro Pro Asn Phe Pro Thr Ala Met Tyr Cys Asp Asn Arg Asn
85 90 95

Leu Lys Tyr Leu Pro Phe Val Pro Ser Arg Met Lys Tyr Val Tyr Phe

100					105					110					
Gln	Asn	Asn	Gln	Ile	Thr	Ser	Ile	Gln	Glu	Gly	Val	Phe	Asp	Asn	Ala
	115						120					125			
Thr	Gly	Leu	Leu	Trp	Ile	Ala	Leu	His	Gly	Asn	Gln	Ile	Thr	Ser	Asp
	130					135						140			
Lys	Val	Gly	Arg	Lys	Val	Phe	Ser	Lys	Leu	Arg	His	Leu	Glu	Arg	Leu
145						150					155				160
Tyr	Leu	Asp	His	Asn	Asn	Leu	Thr	Arg	Met	Pro	Gly	Pro	Leu	Pro	Arg
				165					170					175	
Ser	Leu	Arg	Glu	Leu	His	Leu	Asp	His	Asn	Gln	Ile	Ser	Arg	Val	Pro
			180					185					190		
Asn	Asn	Ala	Leu	Glu	Gly	Leu	Glu	Asn	Leu	Thr	Ala	Leu	Tyr	Leu	Gln
		195					200					205			
His	Asn	Glu	Ile	Gln	Glu	Val	Gly	Ser	Ser	Met	Arg	Gly	Leu	Arg	Ser
	210					215					220				
Leu	Tyr	Leu	Leu	Asp	Leu	Ser	Tyr	Asn	His	Leu	Arg	Lys	Val	Pro	Asp
225						230					235				240
Gly	Leu	Pro	Ser	Ala	Leu	Glu	Gln	Leu	Tyr	Met	Glu	His	Asn	Asn	Val
				245					250					255	
Tyr	Thr	Val	Pro	Asp	Ser	Tyr	Phe	Arg	Gly	Ala	Pro	Lys	Leu	Leu	Tyr
			260					265					270		
Val	Arg	Leu	Ser	His	Asn	Ser	Leu	Thr	Asn	Asn	Gly	Leu	Ala	Ser	Asn
		275					280					285			
Thr	Phe	Asn	Ser	Ser	Ser	Leu	Leu	Glu	Leu	Asp	Leu	Ser	Tyr	Asn	Gln
	290					295					300				
Leu	Gln	Lys	Ile	Pro	Pro	Val	Asn	Thr	Asn	Leu	Glu	Asn	Leu	Tyr	Leu
305						310					315				320
Gln	Gly	Asn	Arg	Ile	Asn	Glu	Phe	Ser	Ile	Ser	Ser	Phe	Cys	Thr	Val
				325					330					335	

Val Asp Val Val Asn Phe Ser Gln Leu Gln Val Val Arg Leu Asp Gly
340 345 350

Asn Glu Met Lys Arg Ser Ala Met Pro Ala Glu Ala Pro Leu Cys Leu
355 360 365

Arg Leu Ala Ser Leu Ile Glu Ile
370 375

<210> 131
<211> 897
<212> PRT
<213> Homo sapiens

<300>
<308> Swissprot/P32927
<309> 1993-10-01
<313> (1)..(897)

<400> 131

Met Val Leu Ala Gln Gly Leu Leu Ser Met Ala Leu Leu Ala Leu Cys
1 5 10 15

Trp Glu Arg Ser Leu Ala Gly Ala Glu Glu Thr Ile Pro Leu Gln Thr
20 25 30

Leu Arg Cys Tyr Asn Asp Tyr Thr Ser His Ile Thr Cys Arg Trp Ala
35 40 45

Asp Thr Gln Asp Ala Gln Arg Leu Val Asn Val Thr Leu Ile Arg Arg
50 55 60

Val Asn Glu Asp Leu Leu Glu Pro Val Ser Cys Asp Leu Ser Asp Asp
65 70 75 80

Met Pro Trp Ser Ala Cys Pro His Pro Arg Cys Val Pro Arg Arg Cys
85 90 95

Val Ile Pro Cys Gln Ser Phe Val Val Thr Asp Val Asp Tyr Phe Ser
100 105 110

Phe Gln Pro Asp Arg Pro Leu Gly Thr Arg Leu Thr Val Thr Leu Thr
115 120 125

Gln His Val Gln Pro Pro Glu Pro Arg Asp Leu Gln Ile Ser Thr Asp
130 135 140

Gln Asp His Phe Leu Leu Thr Trp Ser Val Ala Leu Gly Ser Pro Gln
145 150 155 160

Ser His Trp Leu Ser Pro Gly Asp Leu Glu Phe Glu Val Val Tyr Lys
165 170 175

Arg Leu Gln Asp Ser Trp Glu Asp Ala Ala Ile Leu Leu Ser Asn Thr
180 185 190

Ser Gln Ala Thr Leu Gly Pro Glu His Leu Met Pro Ser Ser Thr Tyr
195 200 205

Val Ala Arg Val Arg Thr Arg Leu Ala Pro Gly Ser Arg Leu Ser Gly
210 215 220

Arg Pro Ser Lys Trp Ser Pro Glu Val Cys Trp Asp Ser Gln Pro Gly
225 230 235 240

Asp Glu Ala Gln Pro Gln Asn Leu Glu Cys Phe Phe Asp Gly Ala Ala
245 250 255

Val Leu Ser Cys Ser Trp Glu Val Arg Lys Glu Val Ala Ser Ser Val
260 265 270

Ser Phe Gly Leu Phe Tyr Lys Pro Ser Pro Asp Ala Gly Glu Glu Glu
275 280 285

Cys Ser Pro Val Leu Arg Glu Gly Leu Gly Ser Leu His Thr Arg His
290 295 300

His Cys Gln Ile Pro Val Pro Asp Pro Ala Thr His Gly Gln Tyr Ile
305 310 315 320

Val Ser Val Gln Pro Arg Arg Ala Glu Lys His Ile Lys Ser Ser Val
325 330 335

Asn Ile Gln Met Ala Pro Pro Ser Leu Asn Val Thr Lys Asp Gly Asp
340 345 350

Ser Tyr Ser Leu Arg Trp Glu Thr Met Lys Met Arg Tyr Glu His Ile
355 360 365

Asp His Thr Phe Glu Ile Gln Tyr Arg Lys Asp Thr Ala Thr Trp Lys
370 375 380

Asp Ser Lys Thr Glu Thr Leu Gln Asn Ala His Ser Met Ala Leu Pro
385 390 395 400

Ala Leu Glu Pro Ser Thr Arg Tyr Trp Ala Arg Val Arg Val Arg Thr
405 410 415

Ser Arg Thr Gly Tyr Asn Gly Ile Trp Ser Glu Trp Ser Glu Ala Arg
420 425 430

Ser Trp Asp Thr Glu Ser Val Leu Pro Met Trp Val Leu Ala Leu Ile
435 440 445

Val Ile Phe Leu Thr Ile Ala Val Leu Leu Ala Leu Arg Phe Cys Gly
450 455 460

Ile Tyr Gly Tyr Arg Leu Arg Arg Lys Trp Glu Glu Lys Ile Pro Asn
465 470 475 480

Pro Ser Lys Ser His Leu Phe Gln Asn Gly Ser Ala Glu Leu Trp Pro
485 490 495

Pro Gly Ser Met Ser Ala Phe Thr Ser Gly Ser Pro Pro His Gln Gly
500 505 510

Pro Trp Gly Ser Arg Phe Pro Glu Leu Glu Gly Val Phe Pro Val Gly
515 520 525

Phe Gly Asp Ser Glu Val Ser Pro Leu Thr Ile Glu Asp Pro Lys His
530 535 540

Val Cys Asp Pro Pro Ser Gly Pro Asp Thr Thr Pro Ala Ala Ser Asp
545 550 555 560

Leu Pro Thr Glu Gln Pro Pro Ser Pro Gln Pro Gly Pro Pro Ala Ala
565 570 575

Ser His Thr Pro Glu Lys Gln Ala Ser Ser Phe Asp Phe Asn Gly Pro

580	585	590
Tyr Leu Gly Pro Pro His Ser Arg Ser Leu Pro Asp Ile Leu Gly Gln 595 600 605		
Pro Glu Pro Pro Gln Glu Gly Gly Ser Gln Lys Ser Pro Pro Pro Gly 610 615 620		
Ser Leu Glu Tyr Leu Cys Leu Pro Ala Gly Gly Gln Val Gln Leu Val 625 630 635 640		
Pro Leu Ala Gln Ala Met Gly Pro Gly Gln Ala Val Glu Val Glu Arg 645 650 655		
Arg Pro Ser Gln Gly Ala Ala Gly Ser Pro Ser Leu Glu Ser Gly Gly 660 665 670		
Gly Pro Ala Pro Pro Ala Leu Gly Pro Arg Val Gly Gly Gln Asp Gln 675 680 685		
Lys Asp Ser Pro Val Ala Ile Pro Met Ser Ser Gly Asp Thr Glu Asp 690 695 700		
Pro Gly Val Ala Ser Gly Tyr Val Ser Ser Ala Asp Leu Val Phe Thr 705 710 715 720		
Pro Asn Ser Gly Ala Ser Ser Val Ser Leu Val Pro Ser Leu Gly Leu 725 730 735		
Pro Ser Asp Gln Thr Pro Ser Leu Cys Pro Gly Leu Ala Ser Gly Pro 740 745 750		
Pro Gly Ala Pro Gly Pro Val Lys Ser Gly Phe Glu Gly Tyr Val Glu 755 760 765		
Leu Pro Pro Ile Glu Gly Arg Ser Pro Arg Ser Pro Arg Asn Asn Pro 770 775 780		
Val Pro Pro Glu Ala Lys Ser Pro Val Leu Asn Pro Gly Glu Arg Pro 785 790 795 800		
Ala Asp Val Ser Pro Thr Ser Pro Gln Pro Glu Gly Leu Leu Val Leu 805 810 815		

Gln Gln Val Gly Asp Tyr Cys Phe Leu Pro Gly Leu Gly Pro Gly Pro
820 825 830

Leu Ser Leu Arg Ser Lys Pro Ser Ser Pro Gly Pro Gly Pro Glu Ile
835 840 845

Lys Asn Leu Asp Gln Ala Phe Gln Val Lys Lys Pro Pro Gly Gln Ala
850 855 860

Val Pro Gln Val Pro Val Ile Gln Leu Phe Lys Ala Leu Lys Gln Gln
865 870 875 880

Asp Tyr Leu Ser Leu Pro Pro Trp Glu Val Asn Lys Pro Gly Glu Val
885 890 895

Cys

<210> 132
<211> 261
<212> PRT
<213> Homo sapiens

<300>
<308> Swissprot/Q60493
<309> 1996-11-01
<313> (1)..(261)

<400> 132

Met Ile Tyr Lys Cys Pro Met Cys Arg Glu Phe Phe Ser Glu Arg Ala
1 5 10 15

Asp Leu Phe Met His Gln Lys Val His Thr Ala Glu Lys Pro His Lys
20 25 30

Cys Asp Lys Cys Asp Lys Gly Phe Phe His Ile Ser Glu Leu His Ile
35 40 45

His Trp Arg Asp His Thr Gly Glu Lys Val Tyr Lys Cys Asp Asp Cys
50 55 60

Gly Lys Asp Phe Ser Thr Thr Thr Lys Leu Asn Arg His Lys Lys Ile
65 70 75 80

His Thr Val Glu Lys Pro Tyr Lys Cys Tyr Glu Cys Gly Lys Ala Phe
85 90 95

Asn Trp Ser Pro His Leu Gln Ile His Met Arg Val His Thr Gly Glu
100 105 110

Lys Pro Tyr Val Cys Ser Glu Cys Gly Arg Gly Phe Ser Asn Ser Ser
115 120 125

Asn Leu Cys Met His Gln Arg Val His Thr Gly Glu Lys Pro Phe Lys
130 135 140

Cys Glu Glu Cys Gly Lys Ala Phe Arg His Thr Ser Ser Leu Cys Met
145 150 155 160

His Gln Arg Val His Thr Gly Glu Lys Pro Tyr Lys Cys Tyr Glu Cys
165 170 175

Gly Lys Ala Phe Ser Gln Ser Ser Ser Leu Cys Ile His Gln Arg Val
180 185 190

His Thr Gly Glu Lys Pro Tyr Arg Cys Cys Gly Cys Gly Lys Ala Phe
195 200 205

Ser Gln Ser Ser Ser Leu Cys Ile His Gln Arg Val His Thr Gly Glu
210 215 220

Lys Pro Phe Lys Cys Asp Glu Cys Gly Lys Ala Phe Ser Gln Ser Thr
225 230 235 240

Ser Leu Cys Ile His Gln Arg Val His Thr Lys Glu Arg Asn His Leu
245 250 255

Lys Ile Ser Val Ile
260

<210> 133
<211> 296
<212> PRT
<213> Homo sapiens

<300>
<308> Swissprot/P04233
<309> 1987-03-20

<313> (1)..(296)

<400> 133

Met His Arg Arg Arg Ser Arg Ser Cys Arg Glu Asp Gln Lys Pro Val
1 5 10 15

Met Asp Asp Gln Arg Asp Leu Ile Ser Asn Asn Glu Gln Leu Pro Met
20 25 30

Leu Gly Arg Arg Pro Gly Ala Pro Glu Ser Lys Cys Ser Arg Gly Ala
35 40 45

Leu Tyr Thr Gly Phe Ser Ile Leu Val Thr Leu Leu Leu Ala Gly Gln
50 55 60

Ala Thr Thr Ala Tyr Phe Leu Tyr Gln Gln Gln Gly Arg Leu Asp Lys
65 70 75 80

Leu Thr Val Thr Ser Gln Asn Leu Gln Leu Glu Asn Leu Arg Met Lys
85 90 95

Leu Pro Lys Pro Pro Lys Pro Val Ser Lys Met Arg Met Ala Thr Pro
100 105 110

Leu Leu Met Gln Ala Leu Pro Met Gly Ala Leu Pro Gln Gly Pro Met
115 120 125

Gln Asn Ala Thr Lys Tyr Gly Asn Met Thr Glu Asp His Val Met His
130 135 140

Leu Leu Gln Asn Ala Asp Pro Leu Lys Val Tyr Pro Pro Leu Lys Gly
145 150 155 160

Ser Phe Pro Glu Asn Leu Arg His Leu Lys Asn Thr Met Glu Thr Ile
165 170 175

Asp Trp Lys Val Phe Glu Ser Trp Met His His Trp Leu Leu Phe Glu
180 185 190

Met Ser Arg His Ser Leu Glu Gln Lys Pro Thr Asp Ala Pro Pro Lys
195 200 205

Val Leu Thr Lys Cys Gln Glu Glu Val Ser His Ile Pro Ala Val His

210 215 220
 Pro Gly Ser Phe Arg Pro Lys Cys Asp Glu Asn Gly Asn Tyr Leu Pro
 225 230 235 240
 Leu Gln Cys Tyr Gly Ser Ile Gly Tyr Cys Trp Cys Val Phe Pro Asn
 245 250 255
 Gly Thr Glu Val Pro Asn Thr Arg Ser Arg Gly His His Asn Cys Ser
 260 265 270
 Glu Ser Leu Glu Leu Glu Asp Pro Ser Ser Gly Leu Gly Val Thr Lys
 275 280 285
 Gln Asp Leu Gly Pro Val Pro Met
 290 295

<210> 134
 <211> 163
 <212> PRT
 <213> Homo sapiens

<300>
 <308> Swissprot/Q99969
 <309> 2000-05-30
 <313> (1)..(163)

<400> 134

Met Arg Arg Leu Leu Ile Pro Leu Ala Leu Trp Leu Gly Ala Val Gly
 1 5 10 15
 Val Gly Val Ala Glu Leu Thr Glu Ala Gln Arg Arg Gly Leu Gln Val
 20 25 30
 Ala Leu Glu Glu Phe His Lys His Pro Pro Val Gln Trp Ala Phe Gln
 35 40 45
 Glu Thr Ser Val Glu Ser Ala Val Asp Thr Pro Phe Pro Ala Gly Ile
 50 55 60
 Phe Val Arg Leu Glu Phe Lys Leu Gln Gln Thr Ser Cys Arg Lys Arg
 65 70 75 80
 Asp Trp Lys Lys Pro Glu Cys Lys Val Arg Pro Asn Gly Arg Lys Arg
 85 90 95

Lys Cys Leu Ala Cys Ile Lys Leu Gly Ser Glu Asp Lys Val Leu Gly
100 105 110

Arg Leu Val His Cys Pro Ile Glu Thr Gln Val Leu Arg Glu Ala Glu
115 120 125

Glu His Gln Glu Thr Gln Cys Leu Arg Val Gln Arg Ala Gly Glu Asp
130 135 140

Pro His Ser Phe Tyr Phe Pro Gly Gln Phe Ala Phe Ser Lys Ala Leu
145 150 155 160

Pro Arg Ser

<210> 135
<211> 2386
<212> PRT
<213> Homo sapiens

<300>
<308> Swissprot/P02751
<309> 1986-07-21
<313> (1)..(2386)

<400> 135

Met Leu Arg Gly Pro Gly Pro Gly Leu Leu Leu Leu Ala Val Gln Cys
1 5 10 15

Leu Gly Thr Ala Val Pro Ser Thr Gly Ala Ser Lys Ser Lys Arg Gln
20 25 30

Ala Gln Gln Met Val Gln Pro Gln Ser Pro Val Ala Val Ser Gln Ser
35 40 45

Lys Pro Gly Cys Tyr Asp Asn Gly Lys His Tyr Gln Ile Asn Gln Gln
50 55 60

Trp Glu Arg Thr Tyr Leu Gly Asn Ala Leu Val Cys Thr Cys Tyr Gly
65 70 75 80

Gly Ser Arg Gly Phe Asn Cys Glu Ser Lys Pro Glu Ala Glu Glu Thr
85 90 95

Cys Phe Asp Lys Tyr Thr Gly Asn Thr Tyr Arg Val Gly Asp Thr Tyr
100 105 110

Glu Arg Pro Lys Asp Ser Met Ile Trp Asp Cys Thr Cys Ile Gly Ala
115 120 125

Gly Arg Gly Arg Ile Ser Cys Thr Ile Ala Asn Arg Cys His Glu Gly
130 135 140

Gly Gln Ser Tyr Lys Ile Gly Asp Thr Trp Arg Arg Pro His Glu Thr
145 150 155 160

Gly Gly Tyr Met Leu Glu Cys Val Cys Leu Gly Asn Gly Lys Gly Glu
165 170 175

Trp Thr Cys Lys Pro Ile Ala Glu Lys Cys Phe Asp His Ala Ala Gly
180 185 190

Thr Ser Tyr Val Val Gly Glu Thr Trp Glu Lys Pro Tyr Gln Gly Trp
195 200 205

Met Met Val Asp Cys Thr Cys Leu Gly Glu Gly Ser Gly Arg Ile Thr
210 215 220

Cys Thr Ser Arg Asn Arg Cys Asn Asp Gln Asp Thr Arg Thr Ser Tyr
225 230 235 240

Arg Ile Gly Asp Thr Trp Ser Lys Lys Asp Asn Arg Gly Asn Leu Leu
245 250 255

Gln Cys Ile Cys Thr Gly Asn Gly Arg Gly Glu Trp Lys Cys Glu Arg
260 265 270

His Thr Ser Val Gln Thr Thr Ser Ser Gly Ser Gly Pro Phe Thr Asp
275 280 285

Val Arg Ala Ala Val Tyr Gln Pro Gln Pro His Pro Gln Pro Pro Pro
290 295 300

Tyr Gly His Cys Val Thr Asp Ser Gly Val Val Tyr Ser Val Gly Met
305 310 315 320

Gln Trp Leu Lys Thr Gln Gly Asn Lys Gln Met Leu Cys Thr Cys Leu
325 330 335

Gly Asn Gly Val Ser Cys Gln Glu Thr Ala Val Thr Gln Thr Tyr Gly
340 345 350

Gly Asn Ser Asn Gly Glu Pro Cys Val Leu Pro Phe Thr Tyr Asn Gly
355 360 365

Arg Thr Phe Tyr Ser Cys Thr Thr Glu Gly Arg Gln Asp Gly His Leu
370 375 380

Trp Cys Ser Thr Thr Ser Asn Tyr Glu Gln Asp Gln Lys Tyr Ser Phe
385 390 395 400

Cys Thr Asp His Thr Val Leu Val Gln Thr Gln Gly Gly Asn Ser Asn
405 410 415

Gly Ala Leu Cys His Phe Pro Phe Leu Tyr Asn Asn His Asn Tyr Thr
420 425 430

Asp Cys Thr Ser Glu Gly Arg Arg Asp Asn Met Lys Trp Cys Gly Thr
435 440 445

Thr Gln Asn Tyr Asp Ala Asp Gln Lys Phe Gly Phe Cys Pro Met Ala
450 455 460

Ala His Glu Glu Ile Cys Thr Thr Asn Glu Gly Val Met Tyr Arg Ile
465 470 475 480

Gly Asp Gln Trp Asp Lys Gln His Asp Met Gly His Met Met Arg Cys
485 490 495

Thr Cys Val Gly Asn Gly Arg Gly Glu Trp Thr Cys Ile Ala Tyr Ser
500 505 510

Gln Leu Arg Asp Gln Cys Ile Val Asp Asp Ile Thr Tyr Asn Val Asn
515 520 525

Asp Thr Phe His Lys Arg His Glu Glu Gly His Met Leu Asn Cys Thr
530 535 540

Cys Phe Gly Gln Gly Arg Gly Arg Trp Lys Cys Asp Pro Val Asp Gln

545		550		555		560									
Cys	Gln	Asp	Ser	Glu	Thr	Gly	Thr	Phe	Tyr	Gln	Ile	Gly	Asp	Ser	Trp
				565					570					575	
Glu	Lys	Tyr	Val	His	Gly	Val	Arg	Tyr	Gln	Cys	Tyr	Cys	Tyr	Gly	Arg
			580					585					590		
Gly	Ile	Gly	Glu	Trp	His	Cys	Gln	Pro	Leu	Gln	Thr	Tyr	Pro	Ser	Ser
		595					600					605			
Ser	Gly	Pro	Val	Glu	Val	Phe	Ile	Thr	Glu	Thr	Pro	Ser	Gln	Pro	Asn
	610					615					620				
Ser	His	Pro	Ile	Gln	Trp	Asn	Ala	Pro	Gln	Pro	Ser	His	Ile	Ser	Lys
625					630				635						640
Tyr	Ile	Leu	Arg	Trp	Arg	Pro	Lys	Asn	Ser	Val	Gly	Arg	Trp	Lys	Glu
				645					650					655	
Ala	Thr	Ile	Pro	Gly	His	Leu	Asn	Ser	Tyr	Thr	Ile	Lys	Gly	Leu	Lys
			660					665					670		
Pro	Gly	Val	Val	Tyr	Glu	Gly	Gln	Leu	Ile	Ser	Ile	Gln	Gln	Tyr	Gly
		675					680					685			
His	Gln	Glu	Val	Thr	Arg	Phe	Asp	Phe	Thr	Thr	Thr	Ser	Thr	Ser	Thr
	690					695					700				
Pro	Val	Thr	Ser	Asn	Thr	Val	Thr	Gly	Glu	Thr	Thr	Pro	Phe	Ser	Pro
705					710					715					720
Leu	Val	Ala	Thr	Ser	Glu	Ser	Val	Thr	Glu	Ile	Thr	Ala	Ser	Ser	Phe
				725					730					735	
Val	Val	Ser	Trp	Val	Ser	Ala	Ser	Asp	Thr	Val	Ser	Gly	Phe	Arg	Val
			740					745					750		
Glu	Tyr	Glu	Leu	Ser	Glu	Glu	Gly	Asp	Glu	Pro	Gln	Tyr	Leu	Asp	Leu
		755					760					765			
Pro	Ser	Thr	Ala	Thr	Ser	Val	Asn	Ile	Pro	Asp	Leu	Leu	Pro	Gly	Arg
						775					780				

Lys Tyr Ile Val Asn Val Tyr Gln Ile Ser Glu Asp Gly Glu Gln Ser
785 790 795 800

Leu Ile Leu Ser Thr Ser Gln Thr Thr Ala Pro Asp Ala Pro Pro Asp
805 810 815

Pro Thr Val Asp Gln Val Asp Asp Thr Ser Ile Val Val Arg Trp Ser
820 825 830

Arg Pro Gln Ala Pro Ile Thr Gly Tyr Arg Ile Val Tyr Ser Pro Ser
835 840 845

Val Glu Gly Ser Ser Thr Glu Leu Asn Leu Pro Glu Thr Ala Asn Ser
850 855 860

Val Thr Leu Ser Asp Leu Gln Pro Gly Val Gln Tyr Asn Ile Thr Ile
865 870 875 880

Tyr Ala Val Glu Glu Asn Gln Glu Ser Thr Pro Val Val Ile Gln Gln
885 890 895

Glu Thr Thr Gly Thr Pro Arg Ser Asp Thr Val Pro Ser Pro Arg Asp
900 905 910

Leu Gln Phe Val Glu Val Thr Asp Val Lys Val Thr Ile Met Trp Thr
915 920 925

Pro Pro Glu Ser Ala Val Thr Gly Tyr Arg Val Asp Val Ile Pro Val
930 935 940

Asn Leu Pro Gly Glu His Gly Gln Arg Leu Pro Ile Ser Arg Asn Thr
945 950 955 960

Phe Ala Glu Val Thr Gly Leu Ser Pro Gly Val Thr Tyr Tyr Phe Lys
965 970 975

Val Phe Ala Val Ser His Gly Arg Glu Ser Lys Pro Leu Thr Ala Gln
980 985 990

Gln Thr Thr Lys Leu Asp Ala Pro Thr Asn Leu Gln Phe Val Asn Glu
995 1000 1005

Thr	Asp	Ser	Thr	Val	Leu	Val	Arg	Trp	Thr	Pro	Pro	Arg	Ala	Gln
1010						1015					1020			
Ile	Thr	Gly	Tyr	Arg	Leu	Thr	Val	Gly	Leu	Thr	Arg	Arg	Gly	Gln
1025						1030					1035			
Pro	Arg	Gln	Tyr	Asn	Val	Gly	Pro	Ser	Val	Ser	Lys	Tyr	Pro	Leu
1040						1045					1050			
Arg	Asn	Leu	Gln	Pro	Ala	Ser	Glu	Tyr	Thr	Val	Ser	Leu	Val	Ala
1055						1060					1065			
Ile	Lys	Gly	Asn	Gln	Glu	Ser	Pro	Lys	Ala	Thr	Gly	Val	Phe	Thr
1070						1075					1080			
Thr	Leu	Gln	Pro	Gly	Ser	Ser	Ile	Pro	Pro	Tyr	Asn	Thr	Glu	Val
1085						1090					1095			
Thr	Glu	Thr	Thr	Ile	Val	Ile	Thr	Trp	Thr	Pro	Ala	Pro	Arg	Ile
1100						1105					1110			
Gly	Phe	Lys	Leu	Gly	Val	Arg	Pro	Ser	Gln	Gly	Gly	Glu	Ala	Pro
1115						1120					1125			
Arg	Glu	Val	Thr	Ser	Asp	Ser	Gly	Ser	Ile	Val	Val	Ser	Gly	Leu
1130						1135					1140			
Thr	Pro	Gly	Val	Glu	Tyr	Val	Tyr	Thr	Ile	Gln	Val	Leu	Arg	Asp
1145						1150					1155			
Gly	Gln	Glu	Arg	Asp	Ala	Pro	Ile	Val	Asn	Lys	Val	Val	Thr	Pro
1160						1165					1170			
Leu	Ser	Pro	Pro	Thr	Asn	Leu	His	Leu	Glu	Ala	Asn	Pro	Asp	Thr
1175						1180					1185			
Gly	Val	Leu	Thr	Val	Ser	Trp	Glu	Arg	Ser	Thr	Thr	Pro	Asp	Ile
1190						1195					1200			
Thr	Gly	Tyr	Arg	Ile	Thr	Thr	Thr	Pro	Thr	Asn	Gly	Gln	Gln	Gly
1205						1210					1215			

Asn Ser	Leu Glu Glu Val Val	His Ala Asp Gln Ser	Ser Cys Thr
1220	1225	1230	
Phe Asp	Asn Leu Ser Pro Gly	Leu Glu Tyr Asn Val	Ser Val Tyr
1235	1240	1245	
Thr Val	Lys Asp Asp Lys Glu	Ser Val Pro Ile Ser	Asp Thr Ile
1250	1255	1260	
Ile Pro	Ala Val Pro Pro Pro	Thr Asp Leu Arg Phe	Thr Asn Ile
1265	1270	1275	
Gly Pro	Asp Thr Met Arg Val	Thr Trp Ala Pro Pro	Pro Ser Ile
1280	1285	1290	
Asp Leu	Thr Asn Phe Leu Val	Arg Tyr Ser Pro Val	Lys Asn Glu
1295	1300	1305	
Glu Asp	Val Ala Glu Leu Ser	Ile Ser Pro Ser Asp	Asn Ala Val
1310	1315	1320	
Val Leu	Thr Asn Leu Leu Pro	Gly Thr Glu Tyr Val	Val Ser Val
1325	1330	1335	
Ser Ser	Val Tyr Glu Gln His	Glu Ser Thr Pro Leu	Arg Gly Arg
1340	1345	1350	
Gln Lys	Thr Gly Leu Asp Ser	Pro Thr Gly Ile Asp	Phe Ser Asp
1355	1360	1365	
Ile Thr	Ala Asn Ser Phe Thr	Val His Trp Ile Ala	Pro Arg Ala
1370	1375	1380	
Thr Ile	Thr Gly Tyr Arg Ile	Arg His His Pro Glu	His Phe Ser
1385	1390	1395	
Gly Arg	Pro Arg Glu Asp Arg	Val Pro His Ser Arg	Asn Ser Ile
1400	1405	1410	
Thr Leu	Thr Asn Leu Thr Pro	Gly Thr Glu Tyr Val	Val Ser Ile
1415	1420	1425	
Val Ala	Leu Asn Gly Arg Glu	Glu Ser Pro Leu Leu	Ile Gly Gln

1430		1435		1440
Gln Ser Thr Val Ser Asp Val Pro Arg Asp Leu Glu Val Val Ala				
1445		1450		1455
Ala Thr Pro Thr Ser Leu Leu Ile Ser Trp Asp Ala Pro Ala Val				
1460		1465		1470
Thr Val Arg Tyr Tyr Arg Ile Thr Tyr Gly Glu Thr Gly Gly Asn				
1475		1480		1485
Ser Pro Val Gln Glu Phe Thr Val Pro Gly Ser Lys Ser Thr Ala				
1490		1495		1500
Thr Ile Ser Gly Leu Lys Pro Gly Val Asp Tyr Thr Ile Thr Val				
1505		1510		1515
Tyr Ala Val Thr Gly Arg Gly Asp Ser Pro Ala Ser Ser Lys Pro				
1520		1525		1530
Ile Ser Ile Asn Tyr Arg Thr Glu Ile Asp Lys Pro Ser Gln Met				
1535		1540		1545
Gln Val Thr Asp Val Gln Asp Asn Ser Ile Ser Val Lys Trp Leu				
1550		1555		1560
Pro Ser Ser Ser Pro Val Thr Gly Tyr Arg Val Thr Thr Thr Pro				
1565		1570		1575
Lys Asn Gly Pro Gly Pro Thr Lys Thr Lys Thr Ala Gly Pro Asp				
1580		1585		1590
Gln Thr Glu Met Thr Ile Glu Gly Leu Gln Pro Thr Val Glu Tyr				
1595		1600		1605
Val Val Ser Val Tyr Ala Gln Asn Pro Ser Gly Glu Ser Gln Pro				
1610		1615		1620
Leu Val Gln Thr Ala Val Thr Asn Ile Asp Arg Pro Lys Gly Leu				
1625		1630		1635
Ala Phe Thr Asp Val Asp Val Asp Ser Ile Lys Ile Ala Trp Glu				
1640		1645		1650

Ser	Pro	Gln	Gly	Gln	Val	Ser	Arg	Tyr	Arg	Val	Thr	Tyr	Ser	Ser
1655						1660					1665			
Pro	Glu	Asp	Gly	Ile	His	Glu	Leu	Phe	Pro	Ala	Pro	Asp	Gly	Glu
1670						1675					1680			
Glu	Asp	Thr	Ala	Glu	Leu	Gln	Gly	Leu	Arg	Pro	Gly	Ser	Glu	Tyr
1685						1690					1695			
Thr	Val	Ser	Val	Val	Ala	Leu	His	Asp	Asp	Met	Glu	Ser	Gln	Pro
1700						1705					1710			
Leu	Ile	Gly	Thr	Gln	Ser	Thr	Ala	Ile	Pro	Ala	Pro	Thr	Asp	Leu
1715						1720					1725			
Lys	Phe	Thr	Gln	Val	Thr	Pro	Thr	Ser	Leu	Ser	Ala	Gln	Trp	Thr
1730						1735					1740			
Pro	Pro	Asn	Val	Gln	Leu	Thr	Gly	Tyr	Arg	Val	Arg	Val	Thr	Pro
1745						1750					1755			
Lys	Glu	Lys	Thr	Gly	Pro	Met	Lys	Glu	Ile	Asn	Leu	Ala	Pro	Asp
1760						1765					1770			
Ser	Ser	Ser	Val	Val	Val	Ser	Gly	Leu	Met	Val	Ala	Thr	Lys	Tyr
1775						1780					1785			
Glu	Val	Ser	Val	Tyr	Ala	Leu	Lys	Asp	Thr	Leu	Thr	Ser	Arg	Pro
1790						1795					1800			
Ala	Gln	Gly	Val	Val	Thr	Thr	Leu	Glu	Asn	Val	Ser	Pro	Pro	Arg
1805						1810					1815			
Arg	Ala	Arg	Val	Thr	Asp	Ala	Thr	Glu	Thr	Thr	Ile	Thr	Ile	Ser
1820						1825					1830			
Trp	Arg	Thr	Lys	Thr	Glu	Thr	Ile	Thr	Gly	Phe	Gln	Val	Asp	Ala
1835						1840					1845			
Val	Pro	Ala	Asn	Gly	Gln	Thr	Pro	Ile	Gln	Arg	Thr	Ile	Lys	Pro
1850						1855					1860			

Asp	Val	Arg	Ser	Tyr	Thr	Ile	Thr	Gly	Leu	Gln	Pro	Gly	Thr	Asp
1865						1870					1875			
Tyr	Lys	Ile	Tyr	Leu	Tyr	Thr	Leu	Asn	Asp	Asn	Ala	Arg	Ser	Ser
1880						1885					1890			
Pro	Val	Val	Ile	Asp	Ala	Ser	Thr	Ala	Ile	Asp	Ala	Pro	Ser	Asn
1895						1900					1905			
Leu	Arg	Phe	Leu	Ala	Thr	Thr	Pro	Asn	Ser	Leu	Leu	Val	Ser	Trp
1910						1915					1920			
Gln	Pro	Pro	Arg	Ala	Arg	Ile	Thr	Gly	Tyr	Ile	Ile	Lys	Tyr	Glu
1925						1930					1935			
Lys	Pro	Gly	Ser	Pro	Pro	Arg	Glu	Val	Val	Pro	Arg	Pro	Arg	Pro
1940						1945					1950			
Gly	Val	Thr	Glu	Ala	Thr	Ile	Thr	Gly	Leu	Glu	Pro	Gly	Thr	Glu
1955						1960					1965			
Tyr	Thr	Ile	Tyr	Val	Ile	Ala	Leu	Lys	Asn	Asn	Gln	Lys	Ser	Glu
1970						1975					1980			
Pro	Leu	Ile	Gly	Arg	Lys	Lys	Thr	Asp	Glu	Leu	Pro	Gln	Leu	Val
1985						1990					1995			
Thr	Leu	Pro	His	Pro	Asn	Leu	His	Gly	Pro	Glu	Ile	Leu	Asp	Val
2000						2005					2010			
Pro	Ser	Thr	Val	Gln	Lys	Thr	Pro	Phe	Val	Thr	His	Pro	Gly	Tyr
2015						2020					2025			
Asp	Thr	Gly	Asn	Gly	Ile	Gln	Leu	Pro	Gly	Thr	Ser	Gly	Gln	Gln
2030						2035					2040			
Pro	Ser	Val	Gly	Gln	Gln	Met	Ile	Phe	Glu	Glu	His	Gly	Phe	Arg
2045						2050					2055			
Arg	Thr	Thr	Pro	Pro	Thr	Thr	Ala	Thr	Pro	Ile	Arg	His	Arg	Pro
2060						2065					2070			

Arg	Pro	Tyr	Pro	Pro	Asn	Val	Gly	Glu	Glu	Ile	Gln	Ile	Gly	His
2075						2080					2085			
Ile	Pro	Arg	Glu	Asp	Val	Asp	Tyr	His	Leu	Tyr	Pro	His	Gly	Pro
2090						2095					2100			
Gly	Leu	Asn	Pro	Asn	Ala	Ser	Thr	Gly	Gln	Glu	Ala	Leu	Ser	Gln
2105						2110					2115			
Thr	Thr	Ile	Ser	Trp	Ala	Pro	Phe	Gln	Asp	Thr	Ser	Glu	Tyr	Ile
2120						2125					2130			
Ile	Ser	Cys	His	Pro	Val	Gly	Thr	Asp	Glu	Glu	Pro	Leu	Gln	Phe
2135						2140					2145			
Arg	Val	Pro	Gly	Thr	Ser	Thr	Ser	Ala	Thr	Leu	Thr	Gly	Leu	Thr
2150						2155					2160			
Arg	Gly	Ala	Thr	Tyr	Asn	Ile	Ile	Val	Glu	Ala	Leu	Lys	Asp	Gln
2165						2170					2175			
Gln	Arg	His	Lys	Val	Arg	Glu	Glu	Val	Val	Thr	Val	Gly	Asn	Ser
2180						2185					2190			
Val	Asn	Glu	Gly	Leu	Asn	Gln	Pro	Thr	Asp	Asp	Ser	Cys	Phe	Asp
2195						2200					2205			
Pro	Tyr	Thr	Val	Ser	His	Tyr	Ala	Val	Gly	Asp	Glu	Trp	Glu	Arg
2210						2215					2220			
Met	Ser	Glu	Ser	Gly	Phe	Lys	Leu	Leu	Cys	Gln	Cys	Leu	Gly	Phe
2225						2230					2235			
Gly	Ser	Gly	His	Phe	Arg	Cys	Asp	Ser	Ser	Arg	Trp	Cys	His	Asp
2240						2245					2250			
Asn	Gly	Val	Asn	Tyr	Lys	Ile	Gly	Glu	Lys	Trp	Asp	Arg	Gln	Gly
2255						2260					2265			
Glu	Asn	Gly	Gln	Met	Met	Ser	Cys	Thr	Cys	Leu	Gly	Asn	Gly	Lys
2270						2275					2280			
Gly	Glu	Phe	Lys	Cys	Asp	Pro	His	Glu	Ala	Thr	Cys	Tyr	Asp	Asp

2285		2290		2295
Gly Lys Thr Tyr His Val	Gly Glu Gln Trp Gln Lys	Glu Tyr Leu		
2300	2305	2310		
Gly Ala Ile Cys Ser Cys Thr	Cys Phe Gly Gly Gln	Arg Gly Trp		
2315	2320	2325		
Arg Cys Asp Asn Cys Arg Arg	Pro Gly Gly Glu Pro	Ser Pro Glu		
2330	2335	2340		
Gly Thr Thr Gly Gln Ser Tyr	Asn Gln Tyr Ser Gln	Arg Tyr His		
2345	2350	2355		
Gln Arg Thr Asn Thr Asn Val	Asn Cys Pro Ile Glu	Cys Phe Met		
2360	2365	2370		
Pro Leu Asp Val Gln Ala Asp	Arg Glu Asp Ser Arg	Glu		
2375	2380	2385		

<210> 136
 <211> 339
 <212> PRT
 <213> Homo sapiens

<300>
 <308> Swissprot/P07858
 <309> 1988-08-01
 <313> (1)..(339)

<400> 136

Met Trp Gln Leu Trp Ala Ser Leu Cys Cys Leu Leu Val Leu Ala Asn
1 5 10 15
Ala Arg Ser Arg Pro Ser Phe His Pro Val Ser Asp Glu Leu Val Asn
20 25 30
Tyr Val Asn Lys Arg Asn Thr Thr Trp Gln Ala Gly His Asn Phe Tyr
35 40 45
Asn Val Asp Met Ser Tyr Leu Lys Arg Leu Cys Gly Thr Phe Leu Gly
50 55 60
Gly Pro Lys Pro Pro Gln Arg Val Met Phe Thr Glu Asp Leu Lys Leu
65 70 75 80

Pro Ala Ser Phe Asp Ala Arg Glu Gln Trp Pro Gln Cys Pro Thr Ile
85 90 95

Lys Glu Ile Arg Asp Gln Gly Ser Cys Gly Ser Cys Trp Ala Phe Gly
100 105 110

Ala Val Glu Ala Ile Ser Asp Arg Ile Cys Ile His Thr Asn Ala His
115 120 125

Val Ser Val Glu Val Ser Ala Glu Asp Leu Leu Thr Cys Cys Gly Ser
130 135 140

Met Cys Gly Asp Gly Cys Asn Gly Gly Tyr Pro Ala Glu Ala Trp Asn
145 150 155 160

Phe Trp Thr Arg Lys Gly Leu Val Ser Gly Gly Leu Tyr Glu Ser His
165 170 175

Val Gly Cys Arg Pro Tyr Ser Ile Pro Pro Cys Glu His His Val Asn
180 185 190

Gly Ser Arg Pro Pro Cys Thr Gly Glu Gly Asp Thr Pro Lys Cys Ser
195 200 205

Lys Ile Cys Glu Pro Gly Tyr Ser Pro Thr Tyr Lys Gln Asp Lys His
210 215 220

Tyr Gly Tyr Asn Ser Tyr Ser Val Ser Asn Ser Glu Lys Asp Ile Met
225 230 235 240

Ala Glu Ile Tyr Lys Asn Gly Pro Val Glu Gly Ala Phe Ser Val Tyr
245 250 255

Ser Asp Phe Leu Leu Tyr Lys Ser Gly Val Tyr Gln His Val Thr Gly
260 265 270

Glu Met Met Gly Gly His Ala Ile Arg Ile Leu Gly Trp Gly Val Glu
275 280 285

Asn Gly Thr Pro Tyr Trp Leu Val Ala Asn Ser Trp Asn Thr Asp Trp
290 295 300

Gly Asp Asn Gly Phe Phe Lys Ile Leu Arg Gly Gln Asp His Cys Gly
 305 310 315 320

Ile Glu Ser Glu Val Val Ala Gly Ile Pro Arg Thr Asp Gln Tyr Trp
 325 330 335

Glu Lys Ile

<210> 137
 <211> 1249
 <212> PRT
 <213> Homo sapiens

<300>
 <308> Swissprot/P29144
 <309> 1992-12-01
 <313> (1)..(1249)

<400> 137

Met Ala Thr Ala Ala Thr Glu Glu Pro Phe Pro Phe His Gly Leu Leu
 1 5 10 15

Pro Lys Lys Glu Thr Gly Ala Ala Ser Phe Leu Cys Arg Tyr Pro Glu
 20 25 30

Tyr Asp Gly Arg Gly Val Leu Ile Ala Val Leu Asp Thr Gly Val Asp
 35 40 45

Pro Gly Ala Pro Gly Met Gln Val Thr Thr Asp Gly Lys Pro Lys Ile
 50 55 60

Val Asp Ile Ile Asp Thr Thr Gly Ser Gly Asp Val Asn Thr Ala Thr
 65 70 75 80

Glu Val Glu Pro Lys Asp Gly Glu Ile Val Gly Leu Ser Gly Arg Val
 85 90 95

Leu Lys Ile Pro Ala Ser Trp Thr Asn Pro Ser Gly Lys Tyr His Ile
 100 105 110

Gly Ile Lys Asn Gly Tyr Asp Phe Tyr Pro Lys Ala Leu Lys Glu Arg
 115 120 125

Ile Gln Lys Glu Arg Lys Glu Lys Ile Trp Asp Pro Val His Arg Val
130 135 140

Ala Leu Ala Glu Ala Cys Arg Lys Gln Glu Glu Phe Asp Val Ala Asn
145 150 155 160

Asn Gly Ser Ser Gln Ala Asn Lys Leu Ile Lys Glu Glu Leu Gln Ser
165 170 175

Gln Val Glu Leu Leu Asn Ser Phe Glu Lys Lys Tyr Ser Asp Pro Gly
180 185 190

Pro Val Tyr Asp Cys Leu Val Trp His Asp Gly Glu Val Trp Arg Ala
195 200 205

Cys Ile Asp Ser Asn Glu Asp Gly Asp Leu Ser Lys Ser Thr Val Leu
210 215 220

Arg Asn Tyr Lys Glu Ala Gln Glu Tyr Gly Ser Phe Gly Thr Ala Glu
225 230 235 240

Met Leu Asn Tyr Ser Val Asn Ile Tyr Asp Asp Gly Asn Leu Leu Ser
245 250 255

Ile Val Thr Ser Gly Gly Ala His Gly Thr His Val Ala Ser Ile Ala
260 265 270

Ala Gly His Phe Pro Glu Glu Pro Glu Arg Asn Gly Val Ala Pro Gly
275 280 285

Ala Gln Ile Leu Ser Ile Lys Ile Gly Asp Thr Arg Leu Ser Thr Met
290 295 300

Glu Thr Gly Thr Gly Leu Ile Arg Ala Met Ile Glu Val Ile Asn His
305 310 315 320

Lys Cys Asp Leu Val Asn Tyr Ser Tyr Gly Glu Ala Thr His Trp Pro
325 330 335

Asn Ser Gly Arg Ile Cys Glu Val Ile Asn Glu Ala Val Trp Lys His
340 345 350

Asn Ile Ile Tyr Val Ser Ser Ala Gly Asn Asn Gly Pro Cys Leu Ser

355							360									365
Thr	Val	Gly	Cys	Pro	Gly	Gly	Thr	Thr	Ser	Ser	Val	Ile	Gly	Val	Gly	
370						375					380					
Ala	Tyr	Val	Ser	Pro	Asp	Met	Met	Val	Ala	Glu	Tyr	Ser	Leu	Arg	Glu	
385					390					395					400	
Lys	Leu	Pro	Ala	Asn	Gln	Tyr	Thr	Trp	Ser	Ser	Arg	Gly	Pro	Ser	Ala	
				405					410					415		
Asp	Gly	Ala	Leu	Gly	Val	Ser	Ile	Ser	Ala	Pro	Gly	Gly	Ala	Ile	Ala	
			420					425					430			
Ser	Val	Pro	Asn	Trp	Thr	Leu	Arg	Gly	Thr	Gln	Leu	Met	Asn	Gly	Thr	
	435						440					445				
Ser	Met	Ser	Ser	Pro	Asn	Ala	Cys	Gly	Gly	Ile	Ala	Leu	Ile	Leu	Ser	
450						455					460					
Gly	Leu	Lys	Ala	Asn	Asn	Ile	Asp	Tyr	Thr	Val	His	Ser	Val	Arg	Arg	
465					470					475					480	
Ala	Leu	Glu	Asn	Thr	Ala	Val	Lys	Ala	Asp	Asn	Ile	Glu	Val	Phe	Ala	
			485						490					495		
Gln	Gly	His	Gly	Ile	Ile	Gln	Val	Asp	Lys	Ala	Tyr	Asp	Tyr	Leu	Val	
			500					505					510			
Gln	Asn	Thr	Ser	Phe	Ala	Asn	Lys	Leu	Gly	Phe	Thr	Val	Thr	Val	Gly	
	515						520					525				
Asn	Asn	Arg	Gly	Ile	Tyr	Leu	Arg	Asp	Pro	Val	Gln	Val	Ala	Ala	Pro	
530						535					540					
Ser	Asp	His	Gly	Val	Gly	Ile	Glu	Pro	Val	Phe	Pro	Glu	Asn	Thr	Glu	
545					550					555					560	
Asn	Ser	Glu	Lys	Ile	Ser	Leu	Gln	Leu	His	Leu	Ala	Leu	Thr	Ser	Asn	
			565					570						575		
Ser	Ser	Trp	Val	Gln	Cys	Pro	Ser	His	Leu	Glu	Leu	Met	Asn	Gln	Cys	
		580					585					590				

Arg His Ile Asn Ile Arg Val Asp Pro Arg Gly Leu Arg Glu Gly Leu
595 600 605

His Tyr Thr Glu Val Cys Gly Tyr Asp Ile Ala Ser Pro Asn Ala Gly
610 615 620

Pro Leu Phe Arg Val Pro Ile Thr Ala Val Ile Ala Ala Lys Val Asn
625 630 635 640

Glu Ser Ser His Tyr Asp Leu Ala Phe Thr Asp Val His Phe Lys Pro
645 650 655

Gly Gln Ile Arg Arg His Phe Ile Glu Val Pro Glu Gly Ala Thr Trp
660 665 670

Ala Glu Val Thr Val Cys Ser Cys Ser Ser Glu Val Ser Ala Lys Phe
675 680 685

Val Leu His Ala Val Gln Leu Val Lys Gln Arg Ala Tyr Arg Ser His
690 695 700

Glu Phe Tyr Lys Phe Cys Ser Leu Pro Glu Lys Gly Thr Leu Thr Glu
705 710 715 720

Ala Phe Pro Val Leu Gly Gly Lys Ala Ile Glu Phe Cys Ile Ala Arg
725 730 735

Trp Trp Ala Ser Leu Ser Asp Val Asn Ile Asp Tyr Thr Ile Ser Phe
740 745 750

His Gly Ile Val Cys Thr Ala Pro Gln Leu Asn Ile His Ala Ser Glu
755 760 765

Gly Ile Asn Arg Phe Asp Val Gln Ser Ser Leu Lys Tyr Glu Asp Leu
770 775 780

Ala Pro Cys Ile Thr Leu Lys Asn Trp Val Gln Thr Leu Arg Pro Val
785 790 795 800

Ser Ala Lys Thr Lys Pro Leu Gly Ser Arg Asp Val Leu Pro Asn Asn
805 810 815

Arg Gln Leu Tyr Glu Met Val Leu Thr Tyr Asn Phe His Gln Pro Lys
820 825 830

Ser Gly Glu Val Thr Pro Ser Cys Pro Leu Leu Cys Glu Leu Leu Tyr
835 840 845

Glu Ser Glu Phe Asp Ser Gln Leu Trp Ile Ile Phe Asp Gln Asn Lys
850 855 860

Arg Gln Met Gly Ser Gly Asp Ala Tyr Pro His Gln Tyr Ser Leu Lys
865 870 875 880

Leu Glu Lys Gly Asp Tyr Thr Ile Arg Leu Gln Ile Arg His Glu Gln
885 890 895

Ile Ser Asp Leu Glu Arg Leu Lys Asp Leu Pro Phe Ile Val Ser His
900 905 910

Arg Leu Ser Asn Thr Leu Ser Leu Asp Ile His Glu Asn His Ser Phe
915 920 925

Ala Leu Leu Gly Lys Lys Lys Ser Ser Asn Leu Thr Leu Pro Pro Lys
930 935 940

Tyr Asn Gln Pro Phe Phe Val Thr Ser Leu Pro Asp Asp Lys Ile Pro
945 950 955 960

Lys Gly Ala Gly Pro Gly Cys Tyr Leu Ala Gly Ser Leu Thr Leu Ser
965 970 975

Lys Thr Glu Leu Gly Lys Lys Ala Asp Val Ile Pro Val His Tyr Tyr
980 985 990

Leu Ile Pro Pro Pro Thr Lys Thr Lys Asn Gly Ser Lys Asp Lys Glu
995 1000 1005

Lys Asp Ser Glu Lys Glu Lys Asp Leu Lys Glu Glu Phe Thr Glu
1010 1015 1020

Ala Leu Arg Asp Leu Lys Ile Gln Trp Met Thr Lys Leu Asp Ser
1025 1030 1035

Ser Asp Ile Tyr Asn Glu Leu Lys Glu Thr Tyr Pro Asn Tyr Leu
1040 1045 1050

Pro Leu Tyr Val Ala Arg Leu His Gln Leu Asp Ala Glu Lys Glu
1055 1060 1065

Arg Met Lys Arg Leu Asn Glu Ile Val Asp Ala Ala Asn Ala Val
1070 1075 1080

Ile Ser His Ile Asp Gln Thr Ala Leu Ala Val Tyr Ile Ala Met
1085 1090 1095

Lys Thr Asp Pro Arg Pro Asp Ala Ala Thr Ile Lys Asn Asp Met
1100 1105 1110

Asp Lys Gln Lys Ser Thr Leu Val Asp Ala Leu Cys Arg Lys Gly
1115 1120 1125

Cys Ala Leu Ala Asp His Leu Leu His Thr Gln Ala Gln Asp Gly
1130 1135 1140

Ala Ile Ser Thr Asp Ala Glu Gly Lys Glu Glu Glu Gly Glu Ser
1145 1150 1155

Pro Leu Asp Ser Leu Ala Glu Thr Phe Trp Glu Thr Thr Lys Trp
1160 1165 1170

Thr Asp Leu Phe Asp Asn Lys Val Leu Thr Phe Ala Tyr Lys His
1175 1180 1185

Ala Leu Val Asn Lys Met Tyr Gly Arg Gly Leu Lys Phe Ala Thr
1190 1195 1200

Lys Leu Val Glu Glu Lys Pro Thr Lys Glu Asn Trp Lys Asn Cys
1205 1210 1215

Ile Gln Leu Met Lys Leu Leu Gly Trp Thr His Cys Ala Ser Phe
1220 1225 1230

Thr Glu Asn Trp Leu Pro Ile Met Tyr Pro Pro Asp Tyr Cys Val
1235 1240 1245

Phe

<210> 138
<211> 433
<212> PRT
<213> Homo sapiens

<300>
<308> Swissprot/Q99538
<309> 1997-11-01
<313> (1)..(433)

<400> 138

Met Val Trp Lys Val Ala Val Phe Leu Ser Val Ala Leu Gly Ile Gly
1 5 10 15

Ala Val Pro Ile Asp Asp Pro Glu Asp Gly Gly Lys His Trp Val Val
20 25 30

Ile Val Ala Gly Ser Asn Gly Trp Tyr Asn Tyr Arg His Gln Ala Asp
35 40 45

Ala Cys His Ala Tyr Gln Ile Ile His Arg Asn Gly Ile Pro Asp Glu
50 55 60

Gln Ile Val Val Met Met Tyr Asp Asp Ile Ala Tyr Ser Glu Asp Asn
65 70 75 80

Pro Thr Pro Gly Ile Val Ile Asn Arg Pro Asn Gly Thr Asp Val Tyr
85 90 95

Gln Gly Val Pro Lys Asp Tyr Thr Gly Glu Asp Val Thr Pro Gln Asn
100 105 110

Phe Leu Ala Val Leu Arg Gly Asp Ala Glu Ala Val Lys Gly Ile Gly
115 120 125

Ser Gly Lys Val Leu Lys Ser Gly Pro Gln Asp His Val Phe Ile Tyr
130 135 140

Phe Thr Asp His Gly Ser Thr Gly Ile Leu Val Phe Pro Asn Glu Asp
145 150 155 160

Leu His Val Lys Asp Leu Asn Glu Thr Ile His Tyr Met Tyr Lys His
165 170 175

Lys Met Tyr Arg Lys Met Val Phe Tyr Ile Glu Ala Cys Glu Ser Gly
180 185 190

Ser Met Met Asn His Leu Pro Asp Asn Ile Asn Val Tyr Ala Thr Thr
195 200 205

Ala Ala Asn Pro Arg Glu Ser Ser Tyr Ala Cys Tyr Tyr Asp Glu Lys
210 215 220

Arg Ser Thr Tyr Leu Gly Asp Trp Tyr Ser Val Asn Trp Met Glu Asp
225 230 235 240

Ser Asp Val Glu Asp Leu Thr Lys Glu Thr Leu His Lys Gln Tyr His
245 250 255

Leu Val Lys Ser His Thr Asn Thr Ser His Val Met Gln Tyr Gly Asn
260 265 270

Lys Thr Ile Ser Thr Met Lys Val Met Gln Phe Gln Gly Met Lys Arg
275 280 285

Lys Ala Ser Ser Pro Val Pro Leu Pro Pro Val Thr His Leu Asp Leu
290 295 300

Thr Pro Ser Pro Asp Val Pro Leu Thr Ile Met Lys Arg Lys Leu Met
305 310 315 320

Asn Thr Asn Asp Leu Glu Glu Ser Arg Gln Leu Thr Glu Glu Ile Gln
325 330 335

Arg His Leu Asp Ala Arg His Leu Ile Glu Lys Ser Val Arg Lys Ile
340 345 350

Val Ser Leu Leu Ala Ala Ser Glu Ala Glu Val Glu Gln Leu Leu Ser
355 360 365

Glu Arg Ala Pro Leu Thr Gly His Ser Cys Tyr Pro Glu Ala Leu Leu
370 375 380

His Phe Arg Thr His Cys Phe Asn Trp His Ser Pro Thr Tyr Glu Tyr
385 390 395 400

Ala Leu Arg His Leu Tyr Val Leu Val Asn Leu Cys Glu Lys Pro Tyr
 405 410 415

Pro Leu His Arg Ile Lys Leu Ser Met Asp His Val Cys Leu Gly His
 420 425 430

Tyr

<210> 139
 <211> 342
 <212> PRT
 <213> Homo sapiens

<300>
 <308> Swissprot/P25105
 <309> 1992-05-01
 <313> (1)..(342)

<400> 139

Met Glu Pro His Asp Ser Ser His Met Asp Ser Glu Phe Arg Tyr Thr
 1 5 10 15

Leu Phe Pro Ile Val Tyr Ser Ile Ile Phe Val Leu Gly Val Ile Ala
 20 25 30

Asn Gly Tyr Val Leu Trp Val Phe Ala Arg Leu Tyr Pro Cys Lys Lys
 35 40 45

Phe Asn Glu Ile Lys Ile Phe Met Val Asn Leu Thr Met Ala Asp Met
 50 55 60

Leu Phe Leu Ile Thr Leu Pro Leu Trp Ile Val Tyr Tyr Gln Asn Gln
 65 70 75 80

Gly Asn Trp Ile Leu Pro Lys Phe Leu Cys Asn Val Ala Gly Cys Leu
 85 90 95

Phe Phe Ile Asn Thr Tyr Cys Ser Val Ala Phe Leu Gly Val Ile Thr
 100 105 110

Tyr Asn Arg Phe Gln Ala Val Thr Arg Pro Ile Lys Thr Ala Gln Ala
 115 120 125

Asn Thr Arg Lys Arg Gly Ile Ser Leu Ser Leu Val Ile Trp Val Ala
130 135 140

Ile Val Gly Ala Ala Ser Tyr Phe Leu Ile Leu Asp Ser Thr Asn Thr
145 150 155 160

Val Pro Asp Ser Ala Gly Ser Gly Asn Val Thr Arg Cys Phe Glu His
165 170 175

Tyr Glu Lys Gly Ser Val Pro Val Leu Ile Ile His Ile Phe Ile Val
180 185 190

Phe Ser Phe Phe Leu Val Phe Leu Ile Ile Leu Phe Cys Asn Leu Val
195 200 205

Ile Ile Arg Thr Leu Leu Met Gln Pro Val Gln Gln Gln Arg Asn Ala
210 215 220

Glu Val Lys Arg Arg Ala Leu Trp Met Val Cys Thr Val Leu Ala Val
225 230 235 240

Phe Ile Ile Cys Phe Val Pro His His Val Val Gln Leu Pro Trp Thr
245 250 255

Leu Ala Glu Leu Gly Phe Gln Asp Ser Lys Phe His Gln Ala Ile Asn
260 265 270

Asp Ala His Gln Val Thr Leu Cys Leu Leu Ser Thr Asn Cys Val Leu
275 280 285

Asp Pro Val Ile Tyr Cys Phe Leu Thr Lys Lys Phe Arg Lys His Leu
290 295 300

Thr Glu Lys Phe Tyr Ser Met Arg Ser Ser Arg Lys Cys Ser Arg Ala
305 310 315 320

Thr Thr Asp Thr Val Thr Glu Val Val Val Pro Phe Asn Gln Ile Pro
325 330 335

Gly Asn Ser Leu Lys Asn
340

<210> 140

<211> 359
<212> PRT
<213> Homo sapiens

<300>
<308> Swissprot/Q92187
<309> 1997-11-01
<313> (1)..(359)

<400> 140

Met Arg Ser Ile Arg Lys Arg Trp Thr Ile Cys Thr Ile Ser Leu Leu
1 5 10 15

Leu Ile Phe Tyr Lys Thr Lys Glu Ile Ala Arg Thr Glu Glu His Gln
20 25 30

Glu Thr Gln Leu Ile Gly Asp Gly Glu Leu Ser Leu Ser Arg Ser Leu
35 40 45

Val Asn Ser Ser Asp Lys Ile Ile Arg Lys Ala Gly Ser Ser Ile Phe
50 55 60

Gln His Asn Val Glu Gly Trp Lys Ile Asn Ser Ser Leu Val Leu Glu
65 70 75 80

Ile Arg Lys Asn Ile Leu Arg Phe Leu Asp Ala Glu Arg Asp Val Ser
85 90 95

Val Val Lys Ser Ser Phe Lys Pro Gly Asp Val Ile His Tyr Val Leu
100 105 110

Asp Arg Arg Arg Thr Leu Asn Ile Ser His Asp Leu His Ser Leu Leu
115 120 125

Pro Glu Val Ser Pro Met Lys Asn Arg Arg Phe Lys Thr Cys Ala Val
130 135 140

Val Gly Asn Ser Gly Ile Leu Leu Asp Ser Glu Cys Gly Lys Glu Ile
145 150 155 160

Asp Ser His Asn Phe Val Ile Arg Cys Asn Leu Ala Pro Val Val Glu
165 170 175

Phe Ala Ala Asp Val Gly Thr Lys Ser Asp Phe Ile Thr Met Asn Pro
180 185 190

Ser Val Val Gln Arg Ala Phe Gly Gly Phe Arg Asn Glu Ser Asp Arg
195 200 205

Glu Lys Phe Val His Arg Leu Ser Met Leu Asn Asp Ser Val Leu Trp
210 215 220

Ile Pro Ala Phe Met Val Lys Gly Gly Glu Lys His Val Glu Trp Val
225 230 235 240

Asn Ala Leu Ile Leu Lys Asn Lys Leu Lys Val Arg Thr Ala Tyr Pro
245 250 255

Ser Leu Arg Leu Ile His Ala Val Arg Gly Tyr Trp Leu Thr Asn Lys
260 265 270

Val Pro Ile Lys Arg Pro Ser Thr Gly Leu Leu Met Tyr Thr Leu Ala
275 280 285

Thr Arg Phe Cys Asp Glu Ile His Leu Tyr Gly Phe Trp Pro Phe Pro
290 295 300

Lys Asp Leu Asn Gly Lys Ala Val Lys Tyr His Tyr Tyr Asp Asp Leu
305 310 315 320

Lys Tyr Arg Tyr Phe Ser Asn Ala Ser Pro His Arg Met Pro Leu Glu
325 330 335

Phe Lys Thr Leu Asn Val Leu His Asn Arg Gly Ala Leu Lys Leu Thr
340 345 350

Thr Gly Lys Cys Val Lys Gln
355

<210> 141
<211> 218
<212> PRT
<213> Homo sapiens

<300>
<308> Swissprot/Q15907
<309> 1998-07-15
<313> (1)..(218)

<400> 141

Met Gly Thr Arg Asp Asp Glu Tyr Asp Tyr Leu Phe Lys Val Val Leu
 1 5 10 15

Ile Gly Asp Ser Gly Val Gly Lys Ser Asn Leu Leu Ser Arg Phe Thr
 20 25 30

Arg Asn Glu Phe Asn Leu Glu Ser Lys Ser Thr Ile Gly Val Glu Phe
 35 40 45

Ala Thr Arg Ser Ile Gln Val Asp Gly Lys Thr Ile Lys Ala Gln Ile
 50 55 60

Trp Asp Thr Ala Gly Gln Glu Arg Tyr Arg Ala Ile Thr Ser Ala Tyr
 65 70 75 80

Tyr Arg Gly Ala Val Gly Ala Leu Leu Val Tyr Asp Ile Ala Lys His
 85 90 95

Leu Thr Tyr Glu Asn Val Glu Arg Trp Leu Lys Glu Leu Arg Asp His
 100 105 110

Ala Asp Ser Asn Ile Val Ile Met Leu Val Gly Asn Lys Ser Asp Leu
 115 120 125

Arg His Leu Arg Ala Val Pro Thr Asp Glu Ala Arg Ala Phe Ala Glu
 130 135 140

Lys Asn Asn Leu Ser Phe Ile Glu Thr Ser Ala Leu Asp Ser Thr Asn
 145 150 155 160

Val Glu Glu Ala Phe Lys Asn Ile Leu Thr Glu Ile Tyr Arg Ile Val
 165 170 175

Ser Gln Lys Gln Ile Ala Asp Arg Ala Ala His Asp Glu Ser Pro Gly
 180 185 190

Asn Asn Val Val Asp Ile Ser Val Pro Pro Thr Thr Asp Gly Gln Lys
 195 200 205

Pro Asn Lys Leu Gln Cys Cys Gln Asn Leu
 210 215

